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                                                                           Sequence 3, Appli
Sequence 47112, A
Sequence 5, Appli
Sequence 30748, A
Sequence 44075, A
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Publication No. US20030087250A1
GENERAL INFORMATION:
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APPLICANT: Manjula GANNAVARAPU
APPLICANT: Sebastian HOERSCH
APPLICANT: Steve G. KOVATS
APPLICANT: Rachel E. MSYERS
APPLICANT: Michael MORRISBY
APPLICANT: Peter OLANDT
APPLICANT: AMI SEN
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Robert C. BAST, Jr.
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Xumei ZHAO
Karen GLATT
   Peter VEIBY
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CURRENT APPLICATION NUMBER: US/10/097,340
CURRENT FILING DATE: 2002-03-14
PRIOR FILING DATE: 2002-03-14
PRIOR FILING DATE: 2001-03-14
PRIOR PLING DATE: 2001-03-14
PRIOR PLING DATE: 2001-03-26
PRIOR PLING DATE: 2001-03-14
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PRIOR FILING DATE: 2001-03-16
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PRIOR FILING DATE: 2001-09-26
PRIOR FILING DATE: 2001-09-19
NUMBER OF SEQ ID NOS: 363
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                                                                      CTCAATGATCCAAAGTTCATTAGTGCCCACCTCATCTCAGAGAGTGACAATCCTGAAGAT
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0y 1880 GGAGACCCACTGACTTACACCATGATAATCACCATGGCCACAGCCT 1939 Db 561 GlyAspProLeuThrHisCysSerAspLeuHisHisAspAshHisHisGlyHisSerPro 580 Oy 1940 GAAGAGAATCATTATGGTAAGAATAGCACATTTTGGAATGCAGTCCGAAG 1999 Db 581 GluGluArgIleIleTyrGlyValGluAsnSerSerThrPheLeuGluCysSerProLys 600 Oy 2000 TCGCAGAGACCTTATGGCAATTCACAGAGAATAGAAAAAAAA	Db 661 LeuLysValThrLeuGluValIleAspThrGluHisLeuGluCluLeuLeuLeuFisLysAsp 680	Oy 2360 ITCIGTGAACAAGTTTGGAAAAGGGACCGAAAACAAGGCCAAAGGCCAGGACATACC 2419 Db 721 PheCysGluGlnValIrpLysArgAspArgLysGlnArgArgGlnArgProGlyHisThr 740 Oy 2420 CCAGGGAACAGTAACAAATGGAAGCACTTACAAGAAAATAAAGAAAG	Qy 2480 ACCCACGAATTTGAGAGGCACCCAGGAGTGTC 2512		K O DEGECK

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                                         1 MetGlyTrpLeuThrArgIleValCySLeuPheTrpGlyValLeuLeuThrAlaArgAla
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Mismatches:
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APPLICATION NUMBER: US/09/060,610
APPLICATION NUMBER: 08/835,268
FILING DATE: *UNKNOWN-
ATTORNEY/AGENT INFORMATION:
NAME: Osman, Richard A.
REGISTRATION NUMBER: 36,627
REFERENCE/DOCKET NUMBER: B94-002-1
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADDRESSEE: SCIENCE & TECHNOLOGY LAW STREET: 268 Bush Street, Suite 3200 CITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Ver
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/067,632
FILING DATE: 04-Feb-2002
CLASSIFICATION: <a href="https://doi.org/10/10/10/10/10/10/">https://doi.org/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10///us/10//us/10///us/10//us/10///us/10///us/10//us/10//us/10//us/10//us/10//us/1
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Matches:
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SEQUENCE CHARACTERISTICS:
LENGTH: 771 amino acids
TYPE: amino acid
TYPE: amino acid
MOLECULE TYPE: protein
SEQUENCE DESCRIPTION: SEQ ID NO
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COMPUTER READABLE FORM:
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Best Local Similarity:
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                                                                                          CysproSerLysThrPheGlyGlyPheAspSerThrLysAspLeuProAspAspVallle 400
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AACTATCAGAATGGGAAGAACAATGTGCCAAGGCTGAAATTATCCTACAAGGAAATGTGG ASSTYCOLAASAGAAATGTGCCCAAGGCTGAAATTATCCTACAAGGAAATGTGG ASSTYCOLAASAGAAATGTGACACACGCCCAAGGAATGTTCATCGTTGGATGGCAACACGCCCAAGAAGAATGTTGATCATTCGTTGGATGGCCAACAGCCCCAAGAAGAATTATCATTCGTTGGATGAAGAAGAAGAACAGCCCAAAAGAATTTTCAATGCTTTCGATGAAGAAAGA	ThrHisAlaArgilediyGinileCysiysAsnAspPheGiyGiyHisArgSerLeuval 280 AATAAATGACAACATTCCTCAAAGCTCGTCTGATTTGCTCAGTGCCAGGTCCAATGGC 1099 ASILYSTPPHATARPHAELEULYSALAArgLeulleCysSerValProGlyProAsnGly 300 ATTGACACTCATTTTGATGAACTTGCAGGATGTATTCCTAATGAACTTTAAAAGATCCTAAA 1159
260 AACTANG 21 ASINTYKI 320 GAATCGG 41 Gluseri 440 GACCTGG 440 GACCTGG 440 GACCTGG 61 LeuLeul 500 GATGAAN 500 GATGAAN 500 GATGAAN 620 TGCACG 620 TGCACG 620 TGCACG 620 TGCACG 630 CATTTGG 630 CATTTGG 640 CATTTGG 650 CATTTGG	261 ThrH 1040 AATA 281 ASIL. 1100 ATTG 301 116A 311 116A 321 ASIL 1220 TGTA 341 CYSM 1280 GGAC 1280 GGAC 1340 TGTC

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741 ProglyAsnSerAsnLysTrpLysHisLeuGlnGluAsnLysLysGlyArgAsnArgArg 760
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                                                                                                                              Sequence 164, Application US/10247671
Publication No. US20030194721A1
GENERAL INFORMATION:
APPLICANT: Mikita, Thomas
APPLICANT: Forter, Gordon, J.
APPLICANT: Forter, Gordon, J.
APPLICANT: Raser, Matthew R.
TITLE OF INVENTION: GENES EXPRESSED IN TREATED FOAM CELLS
FILE REPRENCE: PA-0050 US
CURRENT PILING DATE: 2002-09-18
PRIOR FILING DATE: 2001-09-18
PRIOR FILING DATE: 2001-09-18
NUMBER OF SEQ ID NOS: 186
SOTWARE: PERL PROGram
SEQ ID NO 164
SEQ ID NO 164
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OTHER INFORMATION: Incyte ID No. US20030194721A1 1930967CD1
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Matches:
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                                                   761 ThrHisGluPheGluArgAlaProArgSerVal 771
                                  2480 ACCCACGAATTTGAGAGGCACCCAGGAGTGTC
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ORGANISM: Homo sapiens
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Best Local Similarity:
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; NAME/KEY: misc;
; OTHER INFORMATI
US-10-247-671-164
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Pred. No.:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GGACCCAACTATCAATGGGTGCCTTATCAAGGAAGAGTCCCCTATCCACGGCCAGGAACT 1339
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                301 IleAspThrHisPheAspGluLeuGlnAspValPheLeuMetAsnPheLysAspProLys 320
680 CATITIGAAAACGGCCGTGGGAAGAGICCAIAIGACCCIAAGCTGCTGACAGCAICCCII 739
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                              201 IlePheArgThrLeuGlyKisHisHisProIleArgThrGluGlnHisAspSerArgTrp
                                                                                                                                                181 LeulleAspGlyGluLeuTyrSerGlyThrAlaAlaAspPheMetGlyArgAspPheAla
                                                                                                                                                                                                                                                                                                   860 CTCAATGATCCAAAGTTCATTAGTGCCCACCTCATCTCAGAGAGTGACAATCCTGAAGAT
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                                                                                                                                                                                                 800 ATCTTCCGAACTCTTGGGCACCACCACCCAATCAGGACAGAAGCAGCATGATTCCAGGTGG
                                                                                               740 TTAATAGATGGAGAATTATACTCTGGAACTGCAGCTGATTTTATGGGGCGAGACTTTGCT
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) ORGANISM: Homo sapiens US-10-320-769-3	Alignment Scores: Pred. No.: Score: Score: Score: 655 Percent Similarity: 100.00\$ Best Local Similarity: 100.00\$ Mismatches: Onery Warch Warch 1100.00\$ Mismatches: Onery Warch 1100.00\$	Gaps: (-320-769-3 (1-655)	GCTGGTJ 	Qy 260 AACTATCAGAATGGGAAGAACAATGTGCCAAGGCTGAAATTATCCTACAAGAAATGTTG_319 Dh 21 Benfrund halend and and and and and and and and and a	320 GAATCCAACAATGTGATCACTTTCAATGGCTTGGCCAACAGCTCCAGTTATCATACCTTC [380 CTTTTGGATGAGGAACGGAGTAGGCTGTATGTTGGAGGAACGCATATTTTCATTC (1	440 GACCTGGTTAATATCAAGGATTTTCAAAGATTGTGTGGCCAGTATCTTACACCAGAAGA	81 Aspleuvalasnilelysääspähedinlysilevaltipprovalsertyrthrakgarg 100 500 Gatgaatgcaagtgggctggaaagaacatcctgaaagaatgtgctaatttcatcatcaaggta 55:	ASPGIUCYSLYB'TPALAGIYLYSASPIIELEULYSGIUCYSALAASNPHEIIELYSVAI 12 CTTAAGGCATATAATCAGACTCACTTGTACGCCTGTGGAACGGGGGCTTTTCATCCAATT 61	121 Deubyshialyinshillillilsbeulyinladysstylligiyalarheniskfolie 14 620 TGCACCTACATTGAAATTGGACATCATCTGAGGACAATATTTTTAAGCTGGAGAACTCA 67	141 CysThrTyrIleGlulieGLyHisHisProGluAspAsnilePheLysLeuGluAsnSer 16 680 CATTTGAAAACGGCCGTGGGAAGACTCCATATGACCCTAAGCTGCTGACAGCATCCTT 73	161 HisPheGluAsnGlyArgGlyLysSerProTyrAspProLysLeuLeuThrAlaSerLeu 18 740 TTAATAGATGGAGAATTATACTCTGGAACTGCAGCTGATTTTATGGGGCGAGACTTTGCT 79	DD 181 LeULIEASPGIYGLULEULIYESETGIYINTALAALAASPPNEMETGIYAFGASPPNEALA 200 QY 800 AICTICCGAACTCTIGGGCACCACCACCCCAAICAGGACAGGA	TAGTGCCCACCTCATCTCAGAGAGTGACAATCCTGAAG 	Qy 920 GACAAAGTATACTTTTTTTTTCCGTGAAAATGCAATAGAAAGAA	QY 980 ACTCACGTAGAATAGGTCAGATATGCAAGAATGACTTTGGAGGCACAGAAGTCTGGTG 1039 D 261 ThrHisalaArglleGlyGlnIleCysLysasnAspPheGlyGlyHisargSerLeuVal 280	Qy 1040 AATAAATGGACAACATTCCTCAAAGCTCGTCTGATTTGCTCAGGTCCAAGGTCCAAATGGC 1099 Db 281 ASnLysTrpThrThrPheLeuLysAlaArgLeuIleCysSerValProGlyProAsnGly 300	
Qy 1760 GGGAAAGCGTGTGCCTGAGTCTTGCCTCGCCCGAGACCCTTACTGTGCTTGGGATGCTTCT 1819 Db 521 GIVINSALaCVSALaGIUNNSCVSICALA 1	1820 GOAGACCCACTGACTCTACACTACACATACACACACACACAC	561 GlyAspProLeuThrHisCysSerAspLeuHisHisAspAsnHisHisGlyHisSerPro 580	1940 GAAGAGAATCATCTATGGTGTAGAGAATAGTAGCACATTTTTGGAATGCAGTCGAAG 	OY 2000 TGGCAGAGAGGGGTGGTCTATTGGCAATTCCAGAGGGGAAATGAAGAGGGGAAAAGAAGA 2059	Qy 2060 ATCAGAGTGGATGATCATCATCAGGACAGATCAAGGCCTTCTGCTACGTACG	Qy 2120 CAGAAGGATTCAGGCAATTACCTCTGCCATGGGGTGGAACATGGGTTCATACAAACTCTT 2179	OY .2180 CTTAAGGTAACCCTGGAAGTCATTGACACAGAGCATTTGGAAGAACTTCTTCATAAAGAT 2239 Db 661 LeuLysValThrLeuGluValIleAspThrGluHisLeuGluGluLeuLeuHisLysAsp 680	Qy 2240 GATGATGGAGATGGCTCTAAGACCAAAGAAATGTCCAATAGCATGACACCTAGCCAGAAG 2299	Oy 2300 GTCTGGTACAGACACTTCATGCAGCTCATCAACCCCCAATCTCAACACGATGGATG	Oy 2360 TTCTGTGAACAGTTTGGAAAAGGGCCGAAAACAACGTCGGCAAAGGCCAGGACATGC 2419 Db 721 PheCysGluGlnValTtpLysAtgAspArgLysGlnArgArgGlnArgProGlyHisThr 740	CY 2420 CCAGGGAACAGTAACAATGGAAGCACTTACAAGAAAATAAGAAAGGTAGAAACAGGAGG 2479	Qy 2480 ACCCACGAATTTGAGAGGGCACCCAGGAGTGTC 2512	RESULT 5 US-10-320-769-3 ; Sequence 3, Application US/10320769 ; Publication No. US20030158402A1	; GENERAL INFORMATION: ; APPLICANT: FREEMAN, Gordon J. ; APPLICANT: SCHULTE, Josephin L.	APPLICANT: DOCUSTINS, WESSILAI TITLE OF INVENTION: NUCLEIC ACIDS ENCODING CD100 MOLECULES FILE REPRENCE: DPF-05CPPA2,	CURRENT AFELIATION NUMBER: US/10/320,/09 PRIOR APPLICATION NUMBER: US/08/556,422 PRIOR FILING DATE: 1995-11-09	SET IN SET	

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2060 ATCAGAGTGGATGATCATATCATCAGGACAGATCAAGGCCTTCTGCTACGTAGTCTACAA 2119
               Sequence 47112, Application US/09864761
Patent No. US20020048763A1
GENERAL INFORMATION:
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Score:
Percent Similarity:
151.00
Percent Similarity:
100.00$
Best Local Similarity:
100.00$
Dery Match:
                                                                            APPLICANT: Penn, Sharron G. APPLICANT: Rank, David R. APPLICANT: Hanzel, David K. APPLICANT: Chen, Wensheng
                                                                                                                                                                                                     FILE REFERENCE: Acomica-X-1
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ORGANISM: Homo sapiens
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1100 ATTGACACTCATTTTGATGAACTGCAGGATGTATTCCTAATGAACTTTAAAGATCCTAAA 1159
                                                                              1160 AATCCAGTTGTATATGGAGTGTTTACGACTTCCAGTAACATTTTCAAGGGATCAGCCGTG 1219
                                                                                                                                                                                                                                            1280 GGACCCAACTATCAATGGGTGCCTTATCAAGGAAGAGTCCCCTATCCACGGCCAGGAACT 1339
                                                                                                                                                                                                                                                                                                                           1340 TGTCCCAGCAAAACATTTGGTGGTTTTGACTCCTACAAAGGACCTTCCTGATGATGTTATA 1399
                                                                                                                                                                                                                                                                                                                                                                                                        1400 ACCITIGCAAGAAGICAICCAGCCAIGIACAAICCAGIGIIICCIAIGAACAAICGCCCA 1459
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1460 ATAGTGATCAAAACGGATGTAAATTATCAATTTACACAAATTGTCGTAGACCGAGTGGAT 1519
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1520 GCAGAAAGATGGACAGTATGATGTTATGTTTATCGGAACAGATGTTGGGACCGTTCTTAAA 1579
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1760 GGGAAAGCCTGTGCTGAGTGTTGCCTCGCCGAGACCCTTACTGTGCTTGGGATGGTTCT 1819
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1580 GTAGTTTCAATTCCTAAGGAGACTTGGTATGATTTAGAAGAGGTTCTGCTGGAAGAAATG 1639
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                     301 IleAspThrHisPheAspGluLeuGlnAspValPheLeuMetAsnPheLysAspProLys 320
                                                                                                         321 ASDP. OVALVALTY. GIVVALPHOT THIN THIN SERSER SELITE PHELYSGLY SER JAVA 1340
                                                                                                                                                                                      1940 GAAGAGAGATCATCTATGGTGTAGAGAATAGTAGCACATTTTTGGAATGCAGTCCGAAG 1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2000 TCGCAGAGAGCGCTGGTCTATTGGCAATTCCAGAGGCGAAATGAAGAGGAGAAGAGAAGAGAG 2059
                                                                                                                                                                                                                                                                                                                                                 421 IlevalileLysThrAspValAsnTyrGlnPheThrGlnIleValValAspArgValAsp 440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                441 AlaGluAspGlyGlnTyrAspValMetPhelleGlyThrAspValGlyThrValLeuLys 460
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    481 ThrValPheArgGluProThrAlaIleSerAlaMetGluLeuSerThrLysGlnGlnGln 500
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            621 IleArgValAspAspHisIleIleArgThrAspGlnGlyLeuLeuLeuArgSerLeuGln 640
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TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FOR 
TITLE OF INVENTION: GENE EXPRESSION ANALYSIS BY MICROARRAY
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OTHER INFORMATION: MAP TO AC006322.2

OTHER INFORMATION: EXPRESSED IN LUNG, SIGNAL = 0.73

OTHER INFORMATION: EXPRESSED IN ADULT LIVER, SIGNAL = 0.55

OTHER INFORMATION: SWISSPROT HIT: Q14563, EVALUE 2.00e-88

US-09-864-761-47112
                                                                                                            FILE REFERENCE: 100.05-23
CURRENT APPLICATION NUMBER: US/09/864,761
CURRENT FILING DATE: 2001-05-23
PRIOR APPLICATION NUMBER: US 60/180,312
PRIOR FILING DATE: 2000-05-04
PRIOR FILING DATE: 2000-06-05
PRIOR FILING DATE: 2000-06-05
PRIOR FILING DATE: 2000-06-05
PRIOR FILING DATE: 2000-08-03
PRIOR FILING DATE: 2000-09-04
PRIOR FILING DATE: 2000-09-04
PRIOR FILING DATE: 2000-09-04
PRIOR PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR FILING DATE: 2001-01-30
PRIOR PRIOR DATE: 2001-01-30
PRIOR PRIOR DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
PRIOR PRIOR DATE: 2001
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Mismatches:
Indels:
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Matches:
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RESULT 8

US-10-029-386-30748

US-10-029-386-30748

Sequence 30748, Application US/10029386

Publication No. US20030194704A1

GENERAL INFORMATION:

APPLICANT: Penk, David R.

TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FOR TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXCHANGE USEFUL FOR TITLE OF INVENTION: HUMAN GENOME.*

TITLE OF INVENTION: HUMAN GENOME.*

TITLE OF INVENTION: HUMBER: US/10/029,386

CURRENT APPLICATION NUMBER: US/10/029,386

CURRENT APPLICATION NUMBER: 2001-12-20

NUMBER OF SEQ ID NOS: 34288

SOFTWARE: Annomax Sequence Listing Engine vers. 1.1

SEQ ID NO 30748

LENGTH: 54
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Patent No. US20020048763A1
GENERAL INFORMATION:
APPLICANT: Penn, Sharron G.
APPLICANT: Penn, Sharron G.
APPLICANT: Hanzel, David K.
APPLICANT: Chen, Wensheng
TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FOR TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FOR TITLE OF INVENTION: GENE EXPRESSION ANALYSIS BY MICROARRAY
FILE REFERENCE: Aeonica-X-1
CURRENT APPLICATION NUMBER: US/09/864,761
CURRENT PILING DATE: 2001-05-23
                                                                                        1510
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                  1391 GATGITATAACCTTTGCAAGAAGTCATCCAGCCATGTACAATCCAGTGTTTCCTATGAAC 1450
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              21 AspileTyrGlyLysAlaCysAlaGluCysCysLeuAlaArgAspProTyrCysAlaTrp
                                                                                          1451 AATCGCCCAATAGTGATCAAAACGGATGTAAATTATCAATTTACACAAATTGTCGTAGAC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1691 CAGCAACAACTATATATTGGTTGAACGGCTGGGGTTGCCCAGGTCCCCTTTACACCGGTGT
                                        MAP TO CHR7.1

EXPRESSED IN PLACENTA, SIGNAL = 1.6

EXPRESSED IN FETAL LIVER, SIGNAL = 1.9

EXPRESSED IN LUNG, SIGNAL = 0.59

EXPRESSED IN HERAT, SIGNAL = 2.4

SWISSPROT HIT: Q14563, EVALUE 8.00e-29
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Mismatches:
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Matches:
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OTHER INFORMATION: E
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US-10-029-386-30748
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OTHER INFORMATION:
OTHER INFORMATION:
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Best Local Similarity:
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US-09-773-876-5

Sequence 5, Application US/09773876

Sequence 5, Application US/09973876

Sequence 6, Application US/0993A1

Sequence 7.

Sequence 7.

APPLICANT: Winchester, Robert J.

APPLICANT: Gulko, Percio Saltz

TITLE OF INVENTION: USE OF INIBITORS OF THE ACTIVATION OF CXCR4 RECEPTOR BY SDF-1 IN TITLE OP INVENTION: USE OF INIBITORS OF THE REPRENCE: 0575/57005-A-PCT-US

FILE REFERENCE: 0575/57005-A-PCT-US

CURRENT FILING DATE: 2001-01-31

NUMBER OF SEQ ID NOS: 16

SOFTWARE: Patentin version 3.1
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                                                      2120 CAGAAGGATTCAGGCAATTACCTCTGCCATGCGGTGGAACATGGGTTCATACAAACTCTT 2179
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                                                                                                                                                                                                                                                                                      CCAGGGAACAGTAACAAATGGAAGCACTTACAAGAAAATAAGAAAGGTAGAAACAGGAGG 2479
                                                                                                                                                                                                             GATGATGGAGATGGCTCTAAGACCAAAGAAATGTCCAATAGCATGACACCTAGCCAGAAG 2299
                                                                                                                                                                                                                                                                                                                PheCysGluGlnValTrpLysArgAspArgLysGlnArgArgGlnArgProGlyHisThr 120
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                                                                                    GINLYSASDSerGlyAsnTyrLeuCysHisAlaValGluHisGlyPhelleGlnThrLeu 40
                                                                                                                                                           41 LeulysValThrLeuGluValIleAspThrGluHisLeuGluGluLeuLeuHisLysAsp 60
ProglyThrCysProSerLysThrPheGlyGlyPheAspSerThrLysAspLeuProAsp
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                                                                                                                                   CITAAGGTAACCCTGGAAGTCATTGACACAGAGCATTTGGAAGAACTTCTTCATAAAGAT
                                                                                                                                                                                                                                   61 AspAspGlyAspGlySerLysThrLysGluMetSerAsnSerWetThrProSerGlnLys
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Matches:
Conservative:
Mismatches:
Indels:
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TYPE: PRT
ORGANISM: Homo sapien
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Best Local Similarity:
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US-09-773-876-5
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OTHER INFORMATION: MAP TO CHR7.1

OTHER INFORMATION: EXPRESSED IN ADULT LIVER, SIGNAL = 0.5

OTHER INFORMATION: EXPRESSED IN HELA, SIGNAL = 0.68

OTHER INFORMATION: EXPRESSED IN FETAL LIVER, SIGNAL = 0.64

OTHER INFORMATION: EXPRESSED IN BRAIN, SIGNAL = 0.7

OTHER INFORMATION: EXPRESSED IN BONE MARROW, SIGNAL = 0.69

OTHER INFORMATION: SYPRESSED IN BONE MARROW, SIGNAL = 0.69

OTHER INFORMATION: SYPRESSED IN BONE MARROW, SIGNAL = 0.69
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45 PheSerPheAspLeuValAsnIleLysAspPheGlnLys 57
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CURRENT APPLICATION NUMBER: US/09/864,761
CURRENT FILING DATE: 2001-05-23
PRIOR APPLICATION NUMBER: US 60/180,312
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR PELICATION NUMBER: US 69/632,366
PRIOR APPLICATION NUMBER: US 09/632,366
                                                                                                            Sequence 29372, Application US/10029386 Publication No. US20030194704A1 GENERAL INFORMATION:
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ORGANISM: Homo sapiens
FEATURE:
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Best Local Similarity:
Query Match:
                                                         RESULT 10
US-10-029-386-29372
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Pred. No.:
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SED IN BONE MARROW, SIGNAL = 1.2
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EXPRESSED IN PACENTA, SIGNAL = 1.1
EXPRESSED IN LUNG, SIGNAL = 1.3
EXPRESSED IN FETAL LIVER, SIGNAL = 1.2
EXPRESSED IN FOUT LIVER, SIGNAL = 1.2
EXPRESSED IN BRAIN, SIGNAL = 1.2
EXPRESSED IN BRAIN, SIGNAL = 1.2
SWISSPROT HIT: Q14563, EVALUE 5.00e-25
                        PRIOR FILING DATE: 2000-02-04
PRIOR FILING DATE: 2000-05-26
PRIOR PILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-06-26
PRIOR FILING DATE: 2000-06-03
PRIOR FILING DATE: 2000-06-03
PRIOR FILING DATE: 2000-06-04
PRIOR FILING DATE: 2000-09-07
PRIOR FILING DATE: 2000-09-07
PRIOR PLICATION NUMBER: US 60/236,359
PRIOR PILING DATE: 2001-01-30
PRIOR PELICATION NUMBER: PCT/US01/0066
PRIOR PELICATION NUMBER: PCT/US01/0066
PRIOR PELICATION NUMBER: PCT/US01/0066
PRIOR PELICATION NUMBER: PCT/US01/0066
PRIOR APPLICATION NUMBER: PCT/US01/0066
PRIOR APPLICATION NUMBER: PCT/US01/00669
PRIOR APPLICATION NUMBER: PCT/US01/00670
PRIOR APPLICATION NUMBER: PCT/US01/00670
PRIOR APPLICATION NUMBER: US 09/608,408
PRIOR APPLICATION NUMBER: US 09/774,203
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            US 60/180,312
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ORGANISM: Homo sapiens
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APPLICANT: Penn, Sharron G.
APPLICANT: Rank, David R.
APPLICANT: Rank, David R.
APPLICANT: Hanzel, David R.
TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FOR (TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXCHANGE. ADGUNCA-X-2 CURENT PELLING DATE: 2001-12-20
NUMBER OF SEQ ID NOS: 34288
SEQ ID NO 23372
LENGTH: 60
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Patent No. US20020048763A1

GENERAL INFORMATION:

APPLICANT: Rank, David R.

APPLICANT: Hanzel, David K.

APPLICANT: Chen, Wensheng

TITLE OF INVENTION: GENE EXPRESSION ANALYSIS BY MICROARRAY
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                  APPLICANT: Hanzel, David K.
APPLICANT: Hanzel, David K.
APPLICANT: Hanzel, David K.
TITLE OF INVENTION: GENE EXPRESSION ANALYSIS BY MICROARRAY
FILE REPERENCE: Aconica-X-1
CURRENT APPLICATION NUMBER: US/09/864,761
CURRENT APPLICATION NUMBER: US/09/864,761
CURRENT APPLICATION NUMBER: US/09/864,761
CURRENT FILING DATE: 2001-05-23
REIGN FILING DATE: 2000-02-04
PRICE FILING DATE: 2000-02-04
PRICE FILING DATE: 2000-05-26
PRICE FILING DATE: 2000-05-27
PRICE FILING DATE: 2000-05-27
PRICE FILING DATE: 2001-01-30
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INFORMATION: EXPRESSED IN HEART, SIGNAL = 1.5
INFORMATION: EXPRESSED IN FETAL LIVER, SIGNAL = 1.5
INFORMATION: EXPRESSED IN PLACENTA, SIGNAL = 1.7
INFORMATION: EXPRESSED IN BRAIN, SIGNAL = 1.5
INFORMATION: EXPRESSED IN BARIN, SIGNAL = 1.6
INFORMATION: EXPRESSED IN ADULT LIVER, SIGNAL = 1.6
INFORMATION: EXPRESSED IN ADULT LIVER, SIGNAL = 1.6
INFORMATION: WISSPROT HIT: Q14563, EVALUE 4.000-15
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EXPRESSED IN LUNG, SIGNAL = 0.54
EXPRESSED IN BRAIL SIGNAL = 0.56
EXT HUMAN HIT: BF707780.1, EVALUE 1.00e-17
SWISSPROT HIT: Q14563, EVALUE 1.00e-18
PRIOR APPLICATION NUMBER: GB 24263.6

PRIOR FILING DATE: 2000-10-04

PRIOR PILLING DATE: 2000-10-07

PRIOR PILLING DATE: 2001-03-7

PRIOR PILLING DATE: 2001-01-37

PRIOR FILLING DATE: 2001-01-30

PRIOR APPLICATION NUMBER: PCT/USO1/00664

PRIOR APPLICATION NUMBER: PCT/USO1/00664

PRIOR APPLICATION NUMBER: PCT/USO1/00669

PRIOR PILLING DATE: 2001-01-30

PRIOR PILLING DATE: 2001-01-30

PRIOR PILLING DATE: 2001-01-30

PRIOR APPLICATION NUMBER: PCT/USO1/00665

PRIOR APPLICATION NUMBER: PCT/USO1/00665

PRIOR PILLING DATE: 2001-01-30

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PRIOR FILLING DATE: 2001-01-30

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: PRT
ORGANISM: Homo sapiens
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OTHER INFORMATION: M.
OTHER INFORMATION: E.
OTHER INFORMATION: F.
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Best Local Similarity:
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US-09-864-761-39708
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Query Match
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Length: Matches: Conservative: Mismatches: Indels:

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1766 GCGTGTGCTGAGTGTTGCCTCGCCGAGACCCTTACTGTGCTTGGGATGGT 1816
                                                                                                                                                                           621 AlaCysAlaGluCysCysLeuAlaArgAspProTyrCysAlaTrpAspGly 637
                                                                                                                       US-09-774-490-1 (1-2709) x US-09-813-290-4 (1-782)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-09-774-490-1 (1-2709) x US-09-813-290-2 (1-875)
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                                                                                                                                                                                                                                                          Sequence 2, Application US/09813290
Patent No. US20020042504A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1766 GCGTGTGCTGAGTGTTGCCT
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17.00
100.00%
100.00%
1.90%
4.42e-07
17.00
100.00%
100.00%
1.90%
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ORGANISM: homo sapiens
                                       Percent Similarity:
Best Local Similarity:
Query Match:
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Best Local Similarity:
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APPLICANT: Wilganowski, Nathaniel L.
APPLICANT: Wilganowski, Nathaniel L.
APPLICANT: Wilganowski, Nathaniel L.
APPLICANT: Wilganowski, Nathaniel L.
APPLICANT: Wang, Xiaoming
APPLICANT: Boncho, Gregory
APPLICANT: Boowlile, John
TITLE OF INVENTION: No. US2020042504Alel Human Secreted Proteins and Polymucleotides
TITLE OF INVENTION: No. US2020042504Alel Human Secreted Proteins and Polymucleotides
FILE REFRENCE: LEX-015.USA
CURRENT APPLICATION NUMBER: US 60/190,638
PRIOR APPLICATION NUMBER: US 60/191,188
PRIOR APPLICATION NUMBER: US 60/191,188
PRIOR FILING DATE: 2000-03-22
PRIOR FILING DATE: 2000-03-31
NUMBER OF SEQ ID NOS: 10
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 4
LENGTH: T82
       1766 GCGTGTGCTGAGTGTTGCCTCGCCCGAGACCCTTACTGTGCTTGGGATGGT 1816
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1779
                                                           593 TGTGGAACGGGGCTTTTCATCCAATTTGCACCTACATT 631
                                                                                  CysGlyThrGlyAlaPheHisProlleCysThrTyrlle 33
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Matches:
Conservative:
Mismatches:
Indels:
Gaps:
                                                                                                                                                     Sequence 2, Application US/09731179;
Sequence 2, Application US/09731179;
Patent No. US20010049432A1

GENERAL INFORMATION:
APPLICANT: Holloway, James L..
APPLICANT: Foley, Kevin P.
ITTLE OF INVENTION: HUMAN SEMAPHORIN ZSMF-16
FILE REFERENCE: 99-90;
CURRENT APPLICATION NUMBER: US/09/731,179
CURRENT FILING DATE: 2000-12-06
FRIOR APPLICATION NUMBER: US 60/169,238
PRIOR APPLICATION NUMBER: US 60/169,238
PRIOR FILING DATE: 1999-12-06
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 2
IEDNGTH: 779
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-09-774-490-1 (1-2709) x US-09-731-179-2 (1-779)
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Patent No. US20020004250411
GENERAL INFORMATION:
APPLICANT: Walke, D. Wade
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17.00
100.00$
100.00$
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ORGANISM: Homo Sapiens
US-09-731-179-2
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ORGANISM: homo sapiens
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Query Match:
DB:
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Pred. No.:
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Length:
Matches:
Conservative:
Mismatches:
Indels:

Alignment Scores:

us-09-774-490-1.oligo.rai

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                                                                50, A
Appli
Appli
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Sequence 31, Appl
Sequence 5401, Ap
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Patent No. 5320958
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                7261,
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Sequence 71,
Sequence 726
                                                              Sequence 3
Sequence 2
Sequence 4
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Sequence 2
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Sequence 5
Sequence 5
Sequence 5
                      Sequence Sequence
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MUDIUM TYPE: Floppy disk
COMPUTER: IEM PC compatible
COMPUTER: IEM PC compatible
COMPUTER: IEM PC compatible
COMPUTER: IEM PC-DOS/MS-DOS
SOFTWARE: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/121,713D
FILING DATE: 13-SEP-1993
CLASSIFICATION: 514
US-08-556-422A-6
US-08-556-422A-7
US-08-556-422A-7
US-08-556-422A-7
US-08-556-422A-2
US-08-556-422A-2
US-08-971-089-4
US-09-152-060-58
US-09-252-991A-2158
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US-09-252-991A-2596
US-09-352-91A-2596
US-09-358-296
US-09-358-296
US-09-358-296
US-09-358-296
US-09-369-247-82
US-09-369-247-82
US-09-318-317-418-31
US-08-31976-5363
US-09-31976-5563
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Goodman, Corey S.
APPLICANT: Kolodkin, Alex L.
APPLICANT: Matches, David
APPLICANT: Bentley, David R.
APPLICANT: O'Connor, Timothy
TITLE OF INVENTION: The Semaphorin Gene Family
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE: SCIENCE & TECHNOLOGY LAW GROUP STREET: 268 Bush Street, Suite 3200 CITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ATTORNEY/AGENT INFORMATION:
NAME: Osman, Richard A.
REGISTRATION NUMBER: 36,627
REFERENCE/DOCKET NUMBER: B94-002-1
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 343-4341
TELEFAX: (415) 343-4342
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 1

US-08-121-713D-54

; Sequence 54, Application US/08121713D

; Patent No. 5639856

; GENERAL INFORMATION:
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-MODEL=frame+ n2p.model -DEV=xlp
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-Q=/Cogn2_1/USF09774490/runat_21092004_144321_2615/app_query.fasta_1.2887
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-FGAPEXT=7 -YGAPOP=60 -YGAPEXT=60 -DELEXT=7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /cgn2 6/ptodata/2/jaa/5A COMB.pep:*
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/cgn2 6/ptodata/2/jaa/6A_COMB.pep:*
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/cgn2 6/ptodata/2/jaa/PCTUS COMB.pep:*
/cgn2 6/ptodata/2/jaa/PCTUS COMB.pep:*
                        GenCore version 5.1.6
(c) 1993 - 2004 Compugen Ltd
                                                                                                      protein search, using frame_plus_n2p model
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US-08-835-268-54
US-09-060-610-54
US-09-060-610-54
PCT-US94-10151A-54
US-08-556-42A-3
US-08-556-42A-3
US-08-56-42A-3
US-08-136-42A-3
US-09-311-36-4
US-09-311-461-34
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seq length: 200000000
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Perfect score:
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Maximum 1
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Oy 1040 AATAAATGGACAACATTCCTCAAAGCTCGTCTGATTGCTCAGTGCCAGGTCCAAATGGC 1099 281 ASNLySTrpThrThrPheLeuLySAlaArgLeulleCysSerValProGlyProAsnGly 300 281 ASNLySTrpThrThrPheLeuLySAlaArgLeulleCysSerValProGlyProAsnGly 300 1100 ATTGACACTCATTTTGATGAACTGCAGGATGTATTCCTAATGAACTTTAAAGATCCTAAA 1159 301 IleAspThrHisPheAspGluLeuGlnAspValPheLeuMetAsnPheLySAspProLyS 320 1160 AATCCAGTTGTATATGAGGTGTTTACGACTTCCAGTAACATTTCCAGGGATCAGGGTTCAGGATGTATTACAGGGATCAGGATGTATATGAGGTGTTATATTTTAAGGATAAGATAGAGGTGTTAACAGTAACATGTTAACAATGAACATGTAAGAAGGTGTTCCTTGGTCCATATGCCCAGGGAT 1279 0y 1220 TGTATGTAAAGCATGAGTGAAGAGGGGTTCCTTGGTCCATATGCCCAGGGAT 1279 0y 1220 TGTATGTAAGGATGATGTCAGAAGGGGTTCCTTGGTCCATATGCCCAGGGAT 1279 341 CysMetTyrSerMetSerAspValArgArgValPheLeuGlyProTyrAlaHisArgAsp 350	1280 GGACCCAACTATCAATGGGTGCCTTATCAAGGAAGGTCCCCTATCCACGGCCAGGAACT 133 36. GlyProAsnTytGlnTtpValPtoTytGlnGlyArgValProTytProAtgProGlyThr 380 1340 TGTCCAAGAAACATTTGGTGGTTTTGACTCTACAAAGGACCTTCCTGATGATGTTTATA 139 38. CysProSerLysThrPheGlyGlyPheAspSerThrLysAspLeuProAspAspValle 400	Qy 1400 ACCTTTGCAGAGGCCATGTACAATCCAGGGTTTTCTATGAACAATCGCCCA 1459 Db 401 ThrPheAlaArgSerHisProAlaMetTyrAshProValPheProMetAshAshArgPro 420 Db 401 ThrPheAlaArgSerHisProAlaMetTyrAshProValPheProMetAshAshArgPro 420 Qy 1460 ATAGTGATCAAAGTGAAATTATCAATTTACACAAATTGTCGTAGACCGGATGGAT	1580 GTAGTTTCAAT 461 ValvalSerII 1640 ACAGTTTTTCGG 481 ThrValPheAT 1700 CTAGATTTGG	60 GGGAAAGCGTGTGTGTGTTGCCTCGCCCAGACCCTTACTGCTTGGGATGGTTCT 18 21 GlybyshlacyshlaclucyscysteualaargaspprotyrcysalatrpaspGlyser 54 20 GCATGTTCTGCTATTTTCCACTGCAAAGACGCAAAAACAAAATAAAAAAA 18 41 AlacysSerArgTyrbheProThralalysargArgThrargArgGlnAspIleArgAsn 56 60 GGAACCAACTGACTCAATTCAAACTAAAAAACACATAAACAAAATAASN 56 60 GGAACCAACTGACTCAACTATACAACAATAAAACAAAAAAACAAAC	ProjecthriscysSerAspleuhisHisAspAsnHisHisGlyHisSe GAATCATCTATGGTGAGAATAGTAGCACATTTTGGAATGCAGTCC ASJICISETSTYVALGLASNSerSerThrPheLeuGluCysSerPr GAGCGCTGGTCTATTGGCAATTCCAGAGGCGAAATGAAGGCGAAAAGA HILLINGTHTTTGGCAATTCCAGAGGCGAAATGAAGGGCAAAAGA HYPTTPGLIDHOGGIAATGAAGGCGAATGAAGGGCGAAAGG FIGGATGATCATATCATCAGGACAGATCAAGGCCTTCTGCTACGTAGTGGT FIGGATGATCATATCATCAGGACAGATCAAGGCCTTCTGCTAGTAGTGTGT FIGGATGATCATATCATCAGGACAGATCAAGGCCTTCTGCTACGTAGTGTGT FIGGATGATCATATCATCAGGACAGATCAAGGCCTTCTGCTACGTAGTGTGTGT
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-	TTACTTACAGG 		/alsertyrt) SCTAATTTCA AlaasnPhel 33646CTTTTC 31yAlaPheH	helysleuG TGCTGACAG 	
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ERESTICS: muino acids acid lear protein 771.00 Matches: 100.00% Conservative: 1100.00% Indels: 1 Gaps:	-2709) x US-08-121-713D-54 (1-771) GCTGGTTAACTAGGATTGTCTGTCTTTTCTGGGGAGTA 1yTrpLeuThrArglleValCysLeuFheTrpGlyVal ATCAGAATGGGAAGAACAATGTGCCAAGGCTGAAATTA ATCAGAATGGGAAGAACAATGTGCCAAGGCTGAAATTA	YIGIIMSIIGIYLYSSSIIMSIIGATELONGYSGCUTGCCAACAG CCAACAATGTGATCACTTTCAATGGCTTGGCCAACAG 	euvalasniielysäsppheginlysiievaltrppronarganigation vargementer va	hrTyrileGluileGlyHisHisProGluAspAsnileP TTGAAAACGGCCGTGGGAAGAGTCCATATGACCCTAAGC TTGAAAACGGCCGTGGAAAGAGTCCATATGACCCTAAGC heGluAsnGlyArgGlyLySSerProTyrAspProLySL TAGATGAGAATTATAACTCTGGAACTGCAGCTGATTTTA TAGATGAAATTATAACTCTGAACTGCAAGAGAGGAACTGAACAGAACTGAAACAGAAGAACTGAAACAAGAAGAACAAGAACAAGAACAAGAACAAGAACAAAACAAGAAACAAAAACAAAAACAAGAAACAAAAACAAAAACAAAAACAAAAACAAAAACAAAAACAAAA	heargthreuglyhishishishrolleargthrollud heargthreuglyhishishishrolleargthrollud angartcanagrachtearachashisheulleserglus angarttryttcycogramangcanthagathgan angartythephepheargglumanalaileangan ysvaltyrphephepheargglumanalaileangan accornagantagathagathganagachtrogag accornagantagathagathganagachtrogag isalaargileglydlulecysbyshashagathg
DID NO: 54: BRISTICS: Aid ear protein Comparative: 100.00\$ Mismatches: 100.00\$ Mismatches: 100.00\$ Mismatches: 100.00\$ Mismatches: 100.00\$ Mismatches: 100.00\$	-121-713D-54 ATTGTCTGTCTT 1eValCysLeu AACAATGTGCCA			141 CysThrTyrileGluileGlyHisHisProGluAspAsnIlePheLysLeuGluAsnSer 680 CATTTGAAAACGGCGTGGGAAGAGTCCATATGACCTTAAGCTGCTGACAGCATCCCTT [161 HisPheGluAsnGlyArgGlyLysSerProTyrAspProLysLeuLeuThrAlaSerLeu 740 TTAATAGATGGAGAATTATACTCTGGAACTGCAGCTGATTTATGGGCGGAACTTTTCT 741 [IlePheArgThrLeuGlyHis CTCAATGATCCAAAGTTCATT [

	COSENCINLYS 700 US-09-774-490-1 (1-2709) x US-08-835-268-54 (1-771) COMPARIGACIA STATEMENT OF THE CONTROL OF T	260 AACTATCAGAATGGGAAGAACAATGTGCCAAGGCTGAAATTATCCTACAAAGAATTGTTG 31	Oy 320 GAAICCAACAAIGIGAATCAACTITICAAIGGCITGGCCAACAGCTCCAGTTATCATACCTTC 37:	Oy 380 CTTTTGGATGAGGAAGGGTGTATGTTGGAGGAAAGGATCACATATTTCATTC 43:	Qy 440 GACCTGGTTAATATCAAGGATTTTCAAAAGATTGTGGCCAGTATCTTACACCAGAAGA 499	QY S00 GANGAATGCAAGTGGGCTGGAAAAGACATCCTGAAAGAATTTCATCAAGGTA 559	Qy 560 CITAAGGCATATAATCAGACTCACTTGTACGCCTGTGGAACGGGGGCTTTTCATCCAATT 61	Oy 620 TGCACCTACATTGAACATCGACATCATCCTGAGGACAATATTTTAAGCTGGAGAACTCA 67	Qy 680 CATTTGAAAACGGCGGGGAAGAGTCCATATGACCTAAGCTGCTGACAGCATCCCTT 739	OY 740 TTAATAGAAGAATTATACTCTGGAACTGCAGCTGATTTTATGGGGGGAGACTTTGCT 799	Oy 800 ATCTTCCGAACTCTTGGGCACCACCACCAGGACAGGACA	Oy 860 CTCAATGATCCAAGTTCATTAGTGCCCACCTCATCAGAGAGTGACAATCTCTAAGAGAT 919	Oy 920 GACAAAGTATACTTTTTTTTTTTGGAAAAGGAATAGAAACACTCTGGAAAAGCT 979	Oy 980 ACTCACGCTAGAATAGGTCAGATATGCAAGAATGACTTTGGAGGCCACAGAAGTCTGGTG 103:	1040 AAIAAAIGGACAACATICCICAAACCICICGATITGCICAGGICCAAAIGGC	1100 AITGACACTCATTITGALGAACTGCAGGAIGTATTCCTAATGAACTITAAAGAICCTAAA 11
2120 CAGAAGGATTCAGGCAATTACCTCTGCCATGCGGTGGAACATGGGTTCATACAAACTCTT [681 AspaspdlyAspdlySerLysThrLysGluMetSerAsnSerMetThrProSerGlnLys 2300 GTCTGGTACAGAGACTTCATGCAGCTCATCAACCACCCCAATCTCAACACGATGGATG	2360 TTCTGTGAACAAGTTTGGAAAAGGACCGAAAACAACGTCGGCAAAGGCCAGGACA) 	2420 CCAGGGAACAGTAACAAATGGAAGCACTTACAAGAAAATAAGAAAGGTAGAAACAGGAGG 	2480 ACCCACGAATTIGAGAGGGCACCCAGGAGTGTC 2512 	RESULT 2 US-08-835-268-54 ; Sequence 54, Application US/08835268	Pacent No. 580/826 GENERAL INFORMATION: APPLICANT: Goodman, Corey S. APPLICANT: Kolodkin, Alex L.	9.5.4" HH _H	OF S	CITY: San Francisco STATE: CA COUNTRY: USA ZIP: 94104	COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk COMPUTER: IBP C compatible OPERATING SYSTEM: PC-DOS/MS-DOS	SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/835,268 FILING DATE:	CLASSIFICATION: PRIOR APPLICATION DATA: APPLICATION UNBER: US/08/121,713 FILING DATE: 13-SEP-1993	E 62	TELECOMMUNICATION INFORMATION: TELEPHONE: (415)343-4341 TELEFAX: (415)343-4342 mpr by.	INFORMATION FOR SEQ ID NO: 54: SEQUENCE CHARACTERISTICS: LENGTH: 771 amino acids TYPE: amino acids	GY: li

10 TGTATGTATAGCATGAGTGATGAGAAGGGTGTTCCTTGGTCCATATGCCCACAGGGAT 1	GIVETOASNIYTGINITEVAIETOIYEGINGIYAIGVALFIOIYTETOAGGFTOGIYINE SS TGTCCCAGCAAAACATTTGGTGGTTTTGACTCTACAAAGGACCTTCCTGATGATGTTATA 13 	TCCAGTGT nProvalP TACACAAA		CTGCTGGAAGAATG 1	ACAA 16 nGln 50	CTAINRATIGGITCAACGGCTGGGGTTGCCCGGTTTACAGCGGGTGTGATATITAC 1	GGGAAACGIGIGCTGAGIGTTGCCTCGAGACCCTTACTGICCTGGGATCTTTACTGICCTGGGATCTTTTTTTTTT	GCA1GITCICCCIATITICCCACAGAGACCACAGAGAGAGAGAGAGAGAGAGAGAGA	GGAGGACCACTGACTCACTGTTCAGACTTACACCATGATAATCACCATGGCCACACACCT 19	940 GAAGAGAATCATCTATGGTGTAGAGAATAGTAGCACATTTTTGGAATGCGGGTCGAGG L [106.CARAGAGGCCTGGTCTATIGGCAATICARAGAGGAATIGAGGCGAAATIGAGGCGAAATIGAGGGGAAGAGGAAGAGGAAGAGGAAGAGAGAGA	Alchardidanian chimicalchadanaan chardcolli compositiona chimical chimicalchada chim	GATTCAGGCAATTACCTCTGCCATGGGGTGGAACATGGGTTCATACAACTCTT 217 	%	2240 GATGATGGAGATGGCTCTAAGACCAAAGAAATGTCCAATAGCATGACACCTAGCCAGAAG 2299
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Best Local Similarity: 100.00\$ Mismatches: 0 Query Match: 2 Indels: 0 DB: 2 Gaps: 0 US-09-774-490-1 (1-2709) x US-09-060-692-54 (1-771) QY 200 ATGGGCTGGTTAACTAGGATTGTCTGTCTTTTTTTTTTT	1 METGIJTIPIPIPIPIPIPIPIPIPIPIPIPIPIPIPIPIPIPI	320 GAAICCAACAAIGIGAICACIITCAAIGGCIIGGCCAACAGCICCAGIIAICAIACCIIC 379 	380 CTTTGGATGAGGAACGGAGTAGGCTGTATGTTGGAGCAAAGGATCACATATTTTCATTC 439 [81 ASDLeuValAsnIleLysAspPheGlnLysIleValTrpProValSerTyrThrArgArg 00 GATGAATGCAAGTGGGCTGGAAAAGACATCCTGAAAGAATGTGCTAATTTCATCAAGGA	101 ASPGIUCYSLYSTYPALAGLIYLYSASPILELEULYSGIUCYSALAASBRYNELIELYSVAI 120 560 CTTAAGGCATATAATCAGACTCACTTGTACGCAGGGGGGCTTTTCATCCAATT 619 1		680 CATITIGAAAACGGCCGTGGGAAGAGTCCATATGACCCTAAGCTGCTGACAGCATCCCTT 739 	740 TTAATAGATGGAGAATTATACTCTGGAACTGCAGCTGATTTTATGGGGGGAGACTTTGCT 799 	800 ATCTTCCGAACTCTTGGCACCACCACCCAATCAGGACGAGCAGCATGATTCCAGGTGG 859	860 CTCAATGATCCAAAGTTCATTAGTGCCCACCTCATCTCAGAGAGTGACAATCCTGAAGAT 919 	920 GACAAAGTATACTTTTCTTCCGTGAAAATGCAATAGAGGAACACTCTGGAAAAGCT 979 	980 ACTCACGCTAGAATAGGTCAGATATGCAAGAATGACTTTGGAGGGCACAGAAGTCTGGTG 1039 	AATAAATGGACAACATTCCTCAAAGCTCGTCTGATTTGCTCAGTCCCAGGTCCAAATGGC 1	00 ATTGACACTCATTTTGATGAACTGCAGGATGTATTCCTAATGAACTTTAAAGATCCTAAA	60 AATCCAGTTGTATAIGGAGTTTACGACTTCCAGTAACATTTTCAAGGATCAGCGTG

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        GTCTGGTACAGAGACTTCATGCAGCTCATCAACCACCCCCAATCTCAACACGATGAG
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MEDIUM TYPE: Floppy disk

MEDIUM TYPE: Floppy disk

MEDIUM TYPE: IBM PC compatible

COMPOTER: IBM PC compatible

COMPOTER: IBM PC compatible

COMPUTER: INFORMATION:

APPLICATION NUMBER: US 08/121,713

FILING DATE: 13-SEP-1993

ATTORNEY/AGENT INFORMATION:
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US-08-833-391-54
US-08-833-391-54
| Sequence 54, Application US/08833391
| Patent No. 6013781
| GENERAL INFORMATION:
| APPLICANT: Goodman, Corey S. APPLICANT: Matthes, David APPLICANT: Bentley, David APPLICANT: Bentley, David CAPPLICANT: O'Connor, Timothy TITLE OF INVENTION: The Semaphorin Gene Family CORRESPONDENCES: 100
| CORRESPONDENCES: 100
| CORRESPONDENCE ADDRESS: MOGUND IN COURSES
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Matches:
Conservative:
Mismatches:
Indels:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: Osman, Richard A.
REGISTRATION NUMBER: 36,627
REFERENCE/DOCKET NUMBER: B94-002-1
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 343-4341
TELEFAX: (415) 343-4342
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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Best Local Similarity:
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               AspGluCysLysTrpAlaGlyLysAspIleLeuLysGluCysAlaAsnPheIleLysVal
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CITY: San Francisco
STATE: A COUNTRY: USA
ZIP: 94111-4187
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US94/10151A
FILING DATE: 13-SEP-1994
FILING DATE: 13-SEP-1999
FILING DATE: 13-SEP-199
                                                                                                      Sequence 54, Application PC/TUS9410151A
GENERAL INFORMATION:
APPLICANT: The Regents of the University of California
TITLE OF INVENTION: The Semaphorin Gene Family
NUMBER OF SEQUENCES: 66
CORRESPONDENCE ADDRESS:
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US-08-556-422A-3
Sequence 3, Application US/08556422A
Sequence 3, Application US/08556422A
RESURAL INFORMATION:
APPLICANT: HALL, Kathryn T.
APPLICANT: FREEMAN, Gordon J.
APPLICANT: SCHULTZE, Joachim L.
APPLICANT: SCHULTZE, Joachim L.
APPLICANT: BOUSSIOTIS, Vassiliki
APPLICANT: NADLER, Lee M.
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  MOLECULES
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Matches:
Conservative:
Mismatches:
Indels:
  CD100
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TITLE OF INVENTION: NUCLEIC ACIDS ENCODING CD FILE REFERENCE: DFN-005CPA2, CURRENT APPLICATION NUMBER: US/08/556,422A; CURRENT FILING DATE: 1995-11-09
NUMBER OF SEQ ID NOS: 7
SEQ ID NO 3
SEQ ID NO 3
LENGTH: 655
TYPE: PRT TYPE: PRT TYPE: PRT TYPE: PRT ORGANISM: Homo sapiens
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100.00%
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73.18%
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Best Local Similarity:
Query Match:
DB:
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Pred. No.:
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATE:
FILING DATE: March 11, 1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1778 IGTIGCCTCGCCCGAGACCCTTACTGTGCTTGGGATGGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      529 CysCysLeuAlaArgAspProTyrCysAlaTrpAspGly
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Matches:
Conservative:
Mismatches:
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                                                                                         GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: INAGAKI, Shinobu
APPLICANT: INAGAKI, Shinobu
ITLLE OF INTENTION: NOVEL SERAPHORIN GENES (I)
FILE REFERENCE: 0020-4562P
CURRENT FILING DATE: 1999-05-14
PRIOR APPLICATION NUMBER: ECT/JP97/04111
PRIOR RILING DATE: 1999-11-12
PRIOR PILING DATE: 1999-11-15
NUMBER OF SEQ ID NOS: 4
SOFTWARE: PATCHIN VET: 2.1
SEQ ID NO 1
LENGTH: 775
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE: SCIENCE & TECHNOLOGY LAW GROUP STREET: 75 DENISE DRIVE CITY: HILLSBOROUGH STATE: CALIFORNIA COUNTRY: USA
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APPLICANT: Luo, Yuling
APPLICANT: Xiomei, Xu
TILLE OF INVENTION: Semaphorin K1
NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
                                                                  Sequence 1, Application US/09308179B Patent No. 6436669
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 4, Application US/09041236
Patent No. 6225285
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ATTORNEY/AGENT INFORMATION:
NAME: OSMAN, RICHARD A
REGISTRATION NUMBER: 36,627
REFRENCE/DOCKET NUMBER: EXEI
TELECOMMUNICATION INFORMATION:
TELEPHONE: (650) 343-4341
                                                                                                                                                                                                                                                                                                                                                                                                                                     000703
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INFORMATION FOR SEQ ID NO: 6
SEQUENCE CHARACTERISTICS:
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Best Local Similarit
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Pred. No.:
                                                  US-09-308-179B-1
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                                                                           PheLysAspProLysAsnProValValTyrGlyValPheThrThrSerSerAsn1lePhe 103
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ZIP: 94010

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: JEM PC compatible
TILING DATE: Zem - 2001
CLASSIFICATION: CURKAD A
RECIPIERATION NUMBER: 36,627
REFERENCE/DOCKET NUMBER: EXEL98-001
                                                                                                                                                                                                                                                                                                                                                       Sequence 4, Application US/09771467C

Patent No. 6583277

GENERAL INFORMATION:

TITLE OF INVENTION: Semaphorin K1 Polypeptides

NUMBER OF SEQUENCES: 4

CORRESPONDENCE ADDRESS:

STREET: 75 DENISE DRIVE

CITY: HILLSBOROUGH

STRTE: CALIFORNIA

COUNTRY: USA
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Mismatches:
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Mismatches:
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Matches:
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Matches:
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; SEQUENCE DESCRIPTION: SEQ ID NO: 4:
US-09-771-467C-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELECOMMUNICATION INFORMATION TELEPHONE: (650) 343-434
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 606 amino acids TYPE: amino acid
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INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS
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100.00%
1.12%
: 606 amino acids
amino acid
                                           MOLECULE TYPE: protein
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Best Local Similarity:
Query Match:
DB:
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Query Match:
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Pred. No.:
                 TYPE: ami
TOPOLOGY:
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US-09-771-467C-4
                                                                US-09-041-236-4
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APPLICANT: Hammond, Timothy G.
APPLICANT: Hammond, Timothy G.
APPLICANT: Verroust, Pierre J.
TITLE OF INVENTION: cubilin Protein, DNA Sequences Encoding Cubilin TITLE OF INVENTION: and Uses Thereof FILE REFERENCE: D6148
CURRENT APPLICATION NUMBER: 0200-07-20
PRIOR APPLICATION NUMBER: DCT/US99/01259
PRIOR FILING DATE: 1999-01-21
NUMBER OF SEQ ID NOS: 40
SEQ ID NO 34
LENGTH: 112
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 6, Application US/08556422A
Sequence 6, Application US/08556422A
Patent No. 6576754
GENERAL INFORMATION:
APPLICANT: HALL, Kathryn T.
APPLICANT: SCHULTZE, Joachim L.
APPLICANT: BOUSSIOTIS, Vassiliki
APPLICANT: NDDLER, Lee M.
TITLE OF INVENTION: NUCLEIC ACIDS ENCODING CD100 MOLECULES
FILE REFERENCE: DFN-005CPA.
CURRENT APPLICATION NUMBER: US/08/556,422A
CURRENT FILING DATE: 1995-11-09
NUMBER OF SEQ ID NOS: 7
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255 LysTrpThrThrPheLeuLysAlaArgLeu 264
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                                                                      ; Sequence 34, Application US/09341461; Patent No. 6586389; GENERAL INFORMATION:
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; ORGANISM: Gallus gallus
US-08-556-422A-6
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ORGANISM: human
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US-09-341-461-34
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US-08-556-422A-6
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US-09-341-461-34
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287 LeuAlaArgAspProTyrCysAlaTrp 295

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APPLICANT: HALL, Kathryn T.

APPLICANT: FREEKAN, Gordon J.

APPLICANT: SCHUITZE, Joachim L.

APPLICANT: BOUSSTOTIS, Vassiliki
APPLICANT: NADLER, Lee M.

TITLE OF INVENTION: NUCLEIC ACIDS ENCODING CDIOO MOLECULES

FILE REFERENCE: DFN-005CPA2

CURRENT APPLICATION NUMBER: US/08/556,422A

CURRENT FILING DATE: 1995-11-09

NUMBER OF SEG ID NOS: 7

SOFTWARE: FastSEQ for Windows Version 4.0

SEQ ID NO 4

LENGTH: 607

TYPE: DPM
                                                           APPLICANT: HALL, Kathryn T.
APPLICANT: FREEMAN, Gordon J.
APPLICANT: SCHUTZE, Joachim L.
APPLICANT: BCUSSICIES, Vassiliki
APPLICANT: BCUSSICIES, Vassiliki
APPLICANT: BCUSSICIES, Vassiliki
APPLICANT: NADLER, Lee M.
ITLE OF INVENTION: NUCLEIC ACIDS ENCODING CD100 MOLECULES
FILE REFERENCE: DFN-005CPA2
CURRENT APPLICATION NUMBER: US/08/556,422A
CURRENT FILING DATE: 1995-11-09
NUMBER OF SEQ ID NOS: 7
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 7.
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Matches:
Conservative:
Mismatches:
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Mismatches:
Indels:
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              Sequence 7, Application US/08556422A
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CORGANISM: Homo sapiens
US-08-556-422A-7
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Query Match:
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Query Match:
                                   Patent No. 6576754
GENERAL INFORMATION:
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483 LeuAlaArgAspProTyrCysAlaTrp 491

Search completed: September 21, 2004, 20:35:19 Job time : 97.5 secs

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Sequence 11363, A Sequence 2791, Ap Sequence 11810, A Sequence 28181, A Sequence 2818542, Sequence 2818542, Sequence 2818543, Sequence 2818543, Sequence 2818543, Sequence 10, Appl Sequence 10, Appl Sequence 6, Appli Sequence 7, Appli Sequence 1, Appli Sequence 1, Appli Sequence 1, Appli Sequence 1, Appli Sequence 2, Appli
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Sequence 12826, A
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Sequence 5, Appli
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Bettent No. US20010034332A1
BETTEN No. US20010034332A1
APPLICANT: Jin. Shengfang
TITLE OF INVENTION: RESISTANCE SEQUENCES AND USES THEREOF
FILE REFERENCE: 07334-138001
CURRENT APPLICATION NUMBER: US/09/774,490
CURRENT PILING DATE: 2000.-01-31
PRIOR PILING DATE: 2000-01-31
                  5 US 10-029-386-2791

US-09-864-761-11810

US-09-864-761-6056

US-09-864-761-22893

US-09-864-761-22893

US-09-864-761-22893

US-09-74-490-6

3 US-10-027-632-238542

6 US-10-027-632-238542

6 US-09-908-975-15846

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US-09-897-511A-10

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3 US-08-887-511A-10

3 US-08-887-511A-10

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Pred. No. 0;
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SOFTWARE: FastSEQ for Windows Version 4.0
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Matches 2709; Conservative
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Sequence 283, Appl
Sequence 283, Appl
Sequence 53, Appl
Sequence 30992, A
Sequence 736, App
Sequence 736, App
Sequence 136, App
Sequence 18184, A
Sequence 18184, A
Sequence 27943, A
Sequence 27943, A
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| Cgn2_6/ptodata/1/pubpna/US07_NEW_PUB.seq:*
| Cgn2_6/ptodata/1/pubpna/US06_NEW_PUB.seq:*
| Cgn2_6/ptodata/1/pubpna/US06_PUBCOMB.seq:*
| Cgn2_6/ptodata/1/pubpna/US07_NEW_PUB.seq:*
| Cgn2_6/ptodata/1/pubpna/DS07_NEW_PUB.seq:*
| Cgn2_6/ptodata/1/pubpna/US08_PUBCOMB.seq:*
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| Cgn2_6/ptodata/1/pubpna/US08_PUBCOMB.seq:*
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                           GenCore version 5.1.6
Copyright (c) 1993 - 2004 Compugen Ltd.
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US-10-247-671-83

5 US-10-262-538-9

US-09-922-53

US-09-922-217-30992

US-09-984-761-30992

US-09-984-761-30992

US-09-984-761-30992

US-09-884-761-3094

US-10-025-386-18184

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APPLICANT: OGN WOMENS
APPLICANT: Manjula GANRAVARAPU
APPLICANT: School Seilan HORSEGH
APPLICANT: School Seilan HORSEGH
APPLICANT: School G. KOWATS
APPLICANT: School G. KOWATS
APPLICANT: School B. MEYERS
APPLICANT: Michael WORRISEY
APPLICANT: Michael WORRISEY
APPLICANT: Peter CLANDT
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100.0%; Pred. No. 0;
iive 0; Mismatches
                                                                                                                                                  Sequence 283, Application US/10097340; Publication No. US20030087250Ai; GENERAL INFORMATION:
                                                                                                                                                                                                      APPLICANT: John MONAHAN
APPLICANT: Manjula GANNAVARAPU
APPLICANT: Sebastian HOERSCH
APPLICANT: Steve G. KOVATS
APPLICANT: Steve G. KOVATS
APPLICANT: Rachol E. MEYERS
APPLICANT: Rachol E. MEYERS
APPLICANT: Peter OLANDT
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Best Local Similarity 100.
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CORGANISM: Homo sapiens
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                                                                                                                                #1.25
                                                                                                                              Version
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                                                                                                                                           CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/10/067,632
FILING DATE: 04-F6b-2002
CLASSIFICATION OF CURRENT
APPLICATION NUMBER: US/09/060,610
FILING DATE: CURROWN>
APPLICATION NUMBER: US/09/060,610
FILING DATE: CURROWN>
APPLICATION NUMBER: 08/835,268
FILING DATE: CURROWN>
ATTORNEY/AGENT INFORMATION:
AMME: GENERAL ON NUMBER: 36,627
RESTRENCE/COCKET NUMBER: B94-002-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      92.6%; Score 2508; D
100.0%; Pred. No. 0;
ive 0; Mismatches
                                                                                  COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
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SEQUENCE DESCRIPTION: SEQ ID NO: 53:
                                                                                                                                                                                                                                                                                                                                                                                                                     TELECOMMUNICATION INFORMATION:
TELEPHONE: (415)343-4341
                       ZIP: 94104
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 2601 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEFAX: (415) 343-4342
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   INFORMATION FOR SEQ ID NO: 53: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME/KEY: CDS
LOCATION: 16..2331
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  <Unknown>
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Best Local Similarity 100.
Matches 2508; Conservative
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Kolodkin, Alex L.
Matthes, David
Bentley, David R.
O'Connor, Timothy
TITLE OF INVENTION: The Semaphorin Gene Family
NUMBER OF SEQUENCES: 100
CORRESPONDENCE ADDRESS:
ADDRESSE: SCIENCE & TECHNOLOGY LAW GROUP
STREET: 268 Bush Street, Suite 3200
CITY: San Francisco
STATE: CA
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US-10-067-632-53
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	QY 1873 AAGAANTGGAGACCCACTGACTCACTGTTCAGAC	1749	1993	2053	1929	1989	2233	2293	Qy 2353 GGATGAGTTCTGTGAACAGTTTGGAAAAGGGAC 	2229	2473	2533	2593 2409	Db 2469 CCAATTATAAATTAAATCCATGAGTAACTTTCCTCTCTCT	RESULT 6 US-09-864-761-30992 ; Sequence 30992, Application US/09864761	; FACENT NO. USZUOZUGAS/BASAI ; GRNERAL INFORMATION: ; APPLICANT: Penn, Sharron G. ; APPLICANT: Rank, David R.	APPLICANT: Chen, Wensheng
	549 ATCCCTTTTAATAGATGGAGAATTATACTCTGGAACTGCAGCTGATTTTATGGGGGGGAGA 608 793 CTTTGCTATCTTCCGAACTCTTGGGCACCCCCCAATCAGGACAGAACAGAGCAGATTC 852 609 CTTTGCTAATCTCCGAACTCTTGGGCACCACCACAACAGAACAGAACAGAACAGAACAGAATAGATTC 668	CAGGTGGCTCAATGATCCAAAGTTCATTAGTGCCCACCTCATCTCAGAGAGTGACAATCC 91	913 TGAAGATGACAAAGTATACTTTTCTTCCGTGAAAATGCAATAGATGGAGAACACTCTGG 972 	973 AAAAGCTACTCACGCTAGAATAGGTCAGATATGCAAGAATGACTTTGGAGGGCACAGAAG 1032 	1033 TCTGGTGAATAAATGGACAACATTCCTCAAAGGTCGTCTGATTTGCTCAGTGCCAGGTCC 1092	1093 AAATGGCATTGACACTCATTTTGATGAACTGCAGGATGTATTCCTAATGAACTTTAAAGA 1152 	1153 TCCTAAAATCCAGTTGTATATGCAGTGTTTTACGACTTCCAGTAACATTTTCAAGGGATC 1212 	1213 AGCCGTGTGTATGTATAGCATGAGATGTGAGAAGGGTGTTCCTTGGTCCATATGCCCA 1272 	1273 CAGGGAIGGACCCAACTATCAATGGGTGCCTTATCAAGGAAGAGTCCCCTATCCACGGCC 1332 	aggaacttgtcccagcaaa 	1393 TGTTATAACCTTTGCAAGAAGTCATCCAGCCATGTACAATCCAGTGTTTCCTATGAACAA 1452 	1453 TCGCCCAATAGTGATCAAAACGGATGTAAATTATCAATTTACACAAATTGTGGTGGACG 1512 	1513 AGTGGATGCAGAAGATGGACAGTATGATGTTATCGGAACAGATGTTGGGACGT 1572 	1573 TCTTAAAGTAGTTTCAATTCCTAAGGAGACTTGGTATGATTTAGAAGAGGTTCTGCTGGA 1632 	1633 AGAAATGACAGTTTTCGGGAACCGACTGCTATTTCAGCAATGGAGCTTTCCACTAAGCA 1692 	1693 GCAACAACTATATATTGGTTCAACGGCTGGGGTTGCCCAGCTCCCTTTACACGGGGTGTA 1752 	1753 TATTTACGGGAAAGCGTGTGCTGAGTGTTGCCTCGCCCGAGACCCTTACTGTGCTTGGGA 1812
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CGCCCGAGACCCTTACTGTGCTTGGGA 1628 ACTTACACCATGATAATCACCATGGCCA 1932 AGAATAGTAGCACATTTTTGGAATGCAG 1992 ATTCCAGAGGCGAAATGAAGAGCGAAA 1868 RGACAGATCAAGGCCTTCTGCTACGTAG 2112 CCGAAAACAACGTCGGCAAAGGCCAGG 2412 CCATGCGGTGGACATGGGTTCATACA 2172 CACAGAGCATTTGGAAGAACTTCTTCA 2048 CATCAACCACCCAATCTCAACACGAT 2352 GAGTGTCTGAGCTGCATTACCTCTAGA 2348 ATTCCAGAGGCGAAATGAAGAGCGAAA 2052 AGAAATGTCCAATAGCATGACACCTAG 2292 AGAAATGTCCAATAGCATGACACCTAG 2108 CTTACAAGAAAATAAGAAAGGTAGAAA 2472 GAGTGTCTGAGCTGCATTACCTCTAGA 2532 AACTGGAAAAACAAATGCAATATACAT 2592 AACTGGAAAACAAATGCAATATACAT 2408 AATGGTGGGAAATTCAGCTGAGTTCCA 2652 CACAGAGCATTTGGAAGAACTTCTTCA TAATAGGCTTTTTTT 2700 ||||||||||||||| |TAATAGGCTTTTTTT 2516

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               181 GATGATGGAGATGGCTCTAAGACCAAAGAAATGTCCAATAGCATGACACCTAGCCAGAAG
                                                                                                           2218 GGAAGAACTTCTTCATAAAGATGATGATGGAGATGGCTCTAAGGACCAAAGAAATGTCCAA
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TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND DIAGNOSIS
TITLE OF INVENTION: OF COLON CANCER AND METHODS FOR THEIR USE
FILE REFERENCE: 210121.471C13
CURRENT APPLICATION NUMBER: US/09/922,217
CURRENT APPLICATION NUMBER: US/09/922,217
NUMBER OF SEQ 1D NOS: 1124
SOFTWARE: FASTSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                           2480 ACCCACGAATTTGAGAGGGCACCCAGGAGTGTCTGA 2515
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Pred. No. 5e-152;
0; Mismatches 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Xu, Jiangchun
APPLICANT: Lodes, Michael J.
APPLICANT: Secrist, Heather
APPLICANT: Benson, Darin R.
APPLICANT: Magher: Magheline Joy
APPLICANT: Stolk, John A.
APPLICANT: Wang, Tongtong
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 736, Application US/09922217
Patent No. US20020076414A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LOCATION: 57
OTHER INFORMATION: n = A,T,C or
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Smith, Carole Lynn
King, Gordon E.
Wang, Aijun
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Best Local Similarity 99.7%;
Matches 353; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME/KEY: misc_feature
LOCATION: 57
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LENGTH: 354
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               INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FOR INVENTION: GENE EXPRESSION ANALYSIS BY MICROARRAY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2179
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OTHER INFORMATION: EXPRESSED IN LUNG, SIGNAL = 0.73
OTHER INFORMATION: EXPRESSED IN LUNG, SIGNAL = 0.55
OTHER INFORMATION: EXTRESSED IN AUUT LIVER, SIGNAL = 0.55
OTHER INFORMATION: EST HUMAN HIT: BF700780.1, EVALUE 0.00e+00
OTHER INFORMATION: SWISSPROT HIT: Q14563, EVALUE 2.00e-88
US-09-864-761-30992
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Pred. No. 4.8e-207;
0; Mismatches 1; Indels
TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEUR OF INVENTION: GENE EXPRESSION ANALYSIS BY MICROAL FILE OF INVENTION: GENE EXPRESSION ANALYSIS BY MICROAL CURRENT APPLICATION NUMBER: US/09/864,761

PRIOR APPLICATION NUMBER: US 60/180,312

PRIOR FILING DATE: 2000-02-04

PRIOR FILING DATE: 2000-05-26

PRIOR FILING DATE: 2000-06-05

PRIOR FILING DATE: 2000-06-06

PRIOR FILING DATE: 2000-06-07

PRIOR FILING DATE: 2000-06-07

PRIOR FILING DATE: 2000-06-07

PRIOR FILING DATE: 2000-06-07

PRIOR FILING DATE: 2000-00-07

PRIOR FILING DATE: 2001-01-30

PRIOR FILING DATE: 2000-09-21

PRIOR PRIOR FILING DATE: 2000-09-21

PRIOR PRIOR PRILING DATE: 2000-09-21

PRIOR PRIOR PRILING DATE: 2000-09-30

PRIOR PRIOR PRILING DATE: 2001-01-30

PRIOR PRILING DATE: 2001-01-30

PRIOR PRILING DATE: 2001-01-30

PRIOR PRILING DATE: 2000-09-31

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PRIOR PRILING DATE: 2001-01-30

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Best Local Similarity 99.8°
Matches 455; Conservative
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                                                                                                                                                                                                       Sequence 736, Application US/09833263
; Sequence 736, Application US/09833263
; Patent No. US20020110547A1
; GENERAL INFORMATION:
; APPLICANT: Glapper, Jonathan D.
; APPLICANT: Stolk, John A.
; APPLICANT: Stolk, John A.
; APPLICANT: Madeleine J.
; TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND
; TITLE OF INVENTION: COMPOUNDS FOR COLON CANCER AND METHODS FOR THEIR USE
; TITLE OF INVENTION: LONGOUNDS FOR COLON CANCER AND METHODS FOR THEIR USE
; TITLE OF INVENTION NUMBER: US/09/833,263
; CURRENT FILING DATE: 2001-04-10
; NUMBER OF SEQ ID NOS: 1093
; SEQ ID NOS: 1093
; SEQ ID NO 736
; FORCHMARE: FastSEQ for Windows Version 3.0
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TGGAAAAGGGACCGAAAAAAAACA
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                                      300
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                       241 CAATCTCAACACGATGGATGAGTTCTGTGAACAAGTTTGGAAAGGGACCGAAAACAACG
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                                                                                 2398 TCGGCAAAGGCCAGGACATACCCCAGGGAACAGTAACAAATGGAAGCACTTACA
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ilarity 99.7%; Pred. No. 5e-152;
Conservative 0; Mismatches 1;
    2338 CAATCTCAACACGATGGATGAGTTCTGTGAACAAGTT
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US-10-025-380-736
Sequence 736, Application US/10025380
Sequence 736, Application US/10025380
GENERAL INFORMATION:
APPLICANT: Yu, Jiangchun
APPLICANT: Lodes, Michael J.
APPLICANT: Secrist, Heather
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; LOCATION: (1)....(354)
; OTHER INFORMATION: n = A,T,C or G
US-09-833-263-736
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NAME/KEY: misc_feature
LOCATION: (1)...(354)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: DNA
ORGANISM: Homo sapien
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Matches 353; Conserv
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Sequence 1444...
GENERAL INFORMATION:
APPLICANT: Penn, Sharron G.
APPLICANT: Rank, David R.
APPLICANT: Hanzel, David K.
APPLICANT: Chen, Wesheng
TITLE OF INVENTION: GENDE EXPRESSION ANALYSIS BY MICROARRAY
TITLE OF INVENTION: GENDE EXPRESSION ANALYSIS BY MICROARRAY
FILE REPREBUCE: Acond.ca.~1.
CURRENT APPLICATION NUMBER: US/09/864,761
CURRENT APPLICATION NUMBER: US/09/864,761
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           241 CAATCTCAACACGATGAGTTCTGTGAACAAGTTTGGAAAAGGGACCGAAAACAACG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2278 TAGCATGACCTAGCTAGCCAGAAGGTCTGGTACAGAGACTTCATGCAGCTCATCAACCACC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GGAAGAACTTCTTCATAAAGATGATGATGGAGATGGCTCTAAGACCAAAGAAATGTCCAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 CCTTCTGCTACGTAGGTCTACAACAGAAGGATTCAGGCAATTACCTCCTGCCATGCGGNGGA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2008 CCTTCTGCTACGTGTGTACAACAGGATTCAGGCAATTACCTCTGCCATGCGGTGGA
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APPLICANT: Carter, Darrick
TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND DIAGNOSIS
TITLE OF INVENTION: OF COLON CANCER AND METHODS FOR THEIR USE
FILE REFERENCE: 210121.471C14
CURRENT APPLICATION NUMBER: US/10/025,380
CURRENT FILING DATE: 2001-12-19
NUMBER OF SEQ ID NOS: 1129
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 7.6
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11.2%; Score 303; DB 14; Length 3
Best Local Similarity 99.7%; Pred. No. 5e-152;
Matches 353; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME/KEY: misc_feature
| LOCATION: 57
| OTHER INFORMATION: n = A,T,C or G
US-10-028-380-736
Benson, Darin R.
Meagher, Madeleine Joy
Stolk, John A.
Wang, Tongtong
                                                                                               Jiang, Yuqiu
Smith, Carole L.
King, Gordon E.
King, Ajun
Clapper, Jonathan D.
Skeiky, Yasir A. W.
Fanger, Gary R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: DNA ORGANISM: Homo sapiens
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APPLICANT: Penn, Sharron G.
APPLICANT: Penn, Sharron G.
APPLICANT: Rank, David R.
APPLICANT: Hanzel, David R.
TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FOR CATILE OF INVENTION: EXPRESSION ANALYSIS TWO
TITLE OF INVENTION: EXPRESSION ANALYSIS TWO
TITLE OF INVENTION: EXPRESSION ANALYSIS TWO
CURRENT APPLICATION NUMBER: US/10/029,386
CURRENT APPLICATION NUMBER: US/10/029,386
SURRENT FILING DATE: 2001-12-20
NUMBER OF SEQ ID NOS: 34288
SOFTWARE: Annomax Sequence Listing Engine vers. 1.1
SEQ ID NO 18184
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US-0129-386-4484/C
Sequence 4484, Application US/10029386
Sequence 4484, Application US/10029386
Publication No. US20030194704A1
GENERAL INFORMATION:
APPLICANT: Rank, David R.
APPLICANT: Rank, David R.
APPLICANT: Rank, David R.
TITLE OF INVENTION: EXPRESSION ANALYSIS TWO
STELE REFERENCE: AD001-12-20
UNUMBER OF SEQ ID NOS: 34288
SOUTHARE: Annomax Sequence Listing Engine vers. 1.1
LENGTH: 591
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OTHER INFORMATION: EXPRESSED IN PLACENTA, SIGNAL = 1.6

OTHER INFORMATION: EXPRESSED IN PETAL LIVEK, SIGNAL = 1.9

OTHER INFORMATION: EXPRESSED IN LUNG, SIGNAL = 0.59

OTHER INFORMATION: EXPRESSED IN HEART, SIGNAL = 2.4

OTHER INFORMATION: SAWISSPROT HIT: Q1453, EVALUE 4.00e-29

OTHER INFORMATION: SATISTAND HIT: G1657677.1, EVALUE 9.00e-81

US-10-029-386-18184
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OTHER INFORMATION: EXPRESSED IN PLACENTA, SIGNAL = 1.6

OTHER INFORMATION: EXPRESSED IN FETAL LIVER, SIGNAL = 1

OTHER INFORMATION: EXPRESSED IN LUNG, SIGNAL = 0.59

OTHER INFORMATION: EXPRESSED IN HERRY, SIGNAL = 2.4

OTHER INFORMATION: NT HIT: 9116162409, EVALUE 4.000-85
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100.0%; Pred. No. 1.9e-75;
tive 0; Mismatches 0;
                                                                                       ; Sequence 18184, Application US/10029386; Publication No. US20030194704A1; GENERAL INFORMATION:
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Matches 161; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AGGTCTGGTACAGAGACTTCATGCAGCTCATCAACCACCCCAATCTCAACAC 451
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Pred. No. 4.4e-146;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CTHER INFORMATION: MAP TO AC006322.2
OTHER INFORMATION: EXPRESSED IN LUNG, SIGNAL = 0.73
CTHER INFORMATION: EXPRESSED IN ADULT LIVER, SIGNAL = 0.55
US-09-864-761-14444
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: Annomax Sequence Listing Engine vers. 1.1
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100.0%; Pred. No. 4...
0; Mismatches
PRIOR APPLICATION NUMBER: US 60/180,312
PRIOR FILING DATE: 2000-05-20
PRIOR PLLING DATE: 2000-05-26
PRIOR PLLING DATE: 2000-05-26
PRIOR PLLING DATE: 2000-06-30
PRIOR PLLING DATE: 2000-08-30
PRIOR PLLING DATE: 2000-09-30
PRIOR FILING DATE: 2000-09-27
PRIOR PLLING DATE: 2001-01-30
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Best Local Similarity 100.0
Matches 292, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: DNA
ORGANISM: Homo sapiens
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LENGTH: 496
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MESOUL 10-029-386-16491

Sequence 16491, Application US/10029386

Sequence 16491, Application US/10029386

Publication No. US20030194704A1

GENERAL INFORMATION:
APPLICANT: Penn, Sharron G.
APPLICANT: Penn, Sharron G.
APPLICANT: Hanzel, David K.
TITLE OF INVENTION: EXPRESSION ANALYSIS TWO
TITLE OF INVENTION: EXPRESSION ANALYSIS TWO
TITLE OF INVENTION: EXPRESSION ANALYSIS TWO
CURRENT APPLICATION NUMBER: US/10/029,386
CURRENT APPLICATION NUMBER: US/10/029,386
CURRENT FILING DATE: 2001-12-20
NUMBER OF SEQ ID NOS: 34288
SOFTWARE: Annomax Sequence Listing Engine vers: 1.1
SEQ ID NO 16491
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              429
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        41
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           160 AGAAATGITGGAATCCAACAATGTGATCACTTTCAATGGCTTGGCCAACAGCTCCAGTTA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              370 TCATACCTTCCTTTTGGATGAGGAACGGAGTAGGCTGTATGTTGGAGCAAAGGATCACAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  100 TCATACCTTCCTTTGGATGAGGAACGGGGTAGGCTGTATGTTGGAGGAAAGGATCACAT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       310 AGAAATGTTGGAATCCAACAATGTGATCACTTTCAATGGCTTGGCCAACAGCTCCAGTTA
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INFORMATION: EXPRESSED IN BONE MARROW, SIGNAL = 1.2
INFORMATION: EXPRESSED IN PLACENTRA, SIGNAL = 1.1
INFORMATION: EXPRESSED IN LUNG, SIGNAL = 1.3
INFORMATION: EXPRESSED IN BUTAL LIVER, SIGNAL = 1.2
INFORMATION: EXPRESSED IN ADULT LIVER, SIGNAL = 1.2
INFORMATION: EXPRESSED IN BALLY, SIGNAL = 1.2
INFORMATION: BXPRESSED IN BALLY, SIGNAL = 1.2
INFORMATION: BXF HUMAN HIT: SE694785.1, EVALUE 2.30e+00
INFORMATION: SYLESPROT HIT: Q14563, EVALUE 5.00e-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         5.9%; Score 160; DB 15; Length 191;
100.0%; Pred. No. 6.7e-75;
tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 172;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 EXPRESSED IN ADULT LIVER, SIGNAL = 0.5
EXPRESSED IN HELA, SIGNAL = 0.68
EXPRESSED IN PETAL LIVER, SIGNAL = 0.64
EXPRESSED IN BRAIN, SIGNAL = 0.7
EXPRESSED IN BRAIN, SIGNAL = 0.7
SYRESSED IN BOWE MARROW, SIGNAL = 0.69
SWISSPROT HIT: Q14563, EVALUE 3.00e-25
NI HIT: g15174672, EVALUE 5.00e-85
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100.0%; Pred. No. 6.7e-75;
iive 0; Mismatches 0;
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Best Local Similarity 100.
Matches 160; Conservative
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Best Local Similarity 100.
Matches 160; Conservative
SEQ ID NO 27943
LENGTH: 172
LENGTH: 172
LENGTH: 174
ORGANISM: Home sapiens
FEATURE:
OTHER INFORMATION: EXPRES
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ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FEATURE:
OTHER INFORMATION: MOTHER INFORMATION: EDUTER INFORMATION: EDUTER INFORMATION: EDUTER INFORMATION: EDUTER INFORMATION: EDUTER INFORMATION: STOTHER INFORMATION: STOTHER INFORMATION: STOTHER INFORMATION: N
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OTHER INFORMATIC
US-09-864-761-27943
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APPLICANT: Hanzel, David K.
APPLICANT: Hanzel, David K.
APPLICANT: Hanzel, David K.
TITLE OF INVENTION: HURNA GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FOR TITLE OF INVENTION: HURNA GENOME-DERIVED SINGLE EXPRESSION ARALYSIS BY MICROARRAY FILE REPERBUGE, Ascender X.
CURRENT FAPLICATION: HURNER: US 60/180, 312
PRIOR PELLING DATE: 2001-05-26
PRIOR PELLING DATE: 2000-05-06
PRIOR PELLING DATE: 2000-06-03
PRIOR PELLING DATE: 2001-01-30
PRIOR PELLING DATE: 20
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                                                                                                                                                                                                                                                                                                                  252
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                                                                                                                                                                                                                                                                                                                                                                                                                  251 GATAITTACGGGAAAGCGTGTGCTGAGTGTTGCCTCGCCCGAGACCCTTACTGGCTTGG
                                                                                                                                                                                                                                                                                                                  311 cagcaacaactarararregricaacegecreeegriceccaecreerracaceegrer
                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                 0
   ; CTHER INFORMATION: SWISSPROT HIT: Q14563, EVALUE 2.00e-28; CTHER INFORMATION: EST_HUMAN HIT: BF667677.1, EVALUE 4.00e-80 US-10-029-386-4484
                                                                                                                                    Length 591;
                                                                                                                                                                                                 Indels
                                                                                                                                                                                                                                                              1691 CAGCAACAACTATATTGGTTCAACGGCTGGGGTTGCCCAGCTCCCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GATGGTTCTGCATGTTCTCGCTATTTTCCCACTGCAAAGAG 1851
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: Annomax Sequence Listing Engine vers. 1.1
                                                                                                                                5.9%; Score 161; DB 15;
100.0%; Pred. No. 2.1e-75;
:ive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 27943, Application US/09864761 Patent No. US20020048763A1 GENERAL INFORMATION: SPALICANT: Penn, Sharron G. APPLICANT: Rank, David R. APPLICANT: Hancell, David K. APPLICANT: Chen, Wensheng
                                                                                                                                                                                                       Matches 161; Conservative
                                                                                                                                          Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-09-864-761-27943/c
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Gaps .

Length 484;

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310 AGAAATGTTGGAATCCAACAATGTGATCACTTTCAATGGCTTGGCCAACAGCTCCAGTTA 369
                                                                               370 TCATACCTTCCTTTTGGATGAGGAACGGAGTAGGCTGTATGTTGGAGCAAAAGGATCACAT 429
               Query Match 5.9%; Score 160; DB 9; Length 48
Best Local Similarity 100.0%; Pred. No. 7.3e-75;
Matches 160; Conservative 0; Mismatches 0; Indels
                                                                                                                                                            430 ATTITCATICGACCIGGITAATAICAAGGATTITCAAAAG 469
                                                                                                                                                                                   338 ATTTICATICGACCIGGITAATATCAAGGATTTTCAAAAG 299
                                                                                                                                                                                                                      Search completed: September 23, 2004, 08:32:07
Job time : 2147 secs
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                                                                                                            71 TCATACCTTCCTTTTGGATGAGGAACGGAGTAGGCTGTATGTTGGAGCAAAGGATCACAT 130
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OTHER INFORMATION: EXPRESSED IN BONE MARROW, SIGNAL = 1.2

OTHER INFORMATION: EXPRESSED IN PLACENTA, SIGNAL = 1.1

OTHER INFORMATION: EXPRESSED IN LUNG, SIGNAL = 1.3

OTHER INFORMATION: EXPRESSED IN PETAL LIVER, SIGNAL = 1.1

OTHER INFORMATION: EXPRESSED IN ADULT LIVER, SIGNAL = 1.2

US-09-864-761-11363
                                     430 ATTTTCATTCGACCTGGTTAATATCAAGGATTTTCAAAAG 469
                                                   131 ATTTTCATTCGACCTGGTTAATATCAAGGATTTTCAAAG 170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: Annomax Sequence Listing Engine vers. 1.1
SEQ ID NO 11363
LENGTH: 484
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ORGANISM: Homo sapiens
FEATURE:
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Sequence S

Sequence 1 Sequence 1 Sequence 2 Sequence 3 Sequence 4

Sequence 30, Sequence 7, A Sequence 7, A

Sequence Seguence

Sequence

score:

Title: Perfect :

Sequence:

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Scoring table:

Word size :

Searched:

Database :

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Length 2601;
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CITY: San Francisco
STATE: CA
COUNTRY: USA
ZIP: 94104
COMPUTER READABLE FORM:
MEDIUW TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
COMPUTER: PATENTIN PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/121,713D
FILING DATE: 13-SEP-1993
CLASSIFCATION: S14
ATTORNEY/AGENT INFORMATION:
NAME: OSMEN, Richard A.
REGISTRATION NUMBER: 36,627
REGISTRATION NUMBER: 34,627
RETERBROC/DOCKET NUMBER: B94-002-1
TELERDAM: (415) 343-4341
                                                                                                                                                                                                                                                                                                                                                                                                       Patent No. 5639866
GENERAL INFORMATION:
APPLICANT: Goodman, Corey S.
APPLICANT: Goodman, Corey S.
APPLICANT: Matthes, David R.
APPLICANT: Matthes, David R.
APPLICANT: Matthes, David R.
APPLICANT: Dentley, David R.
APPLICANT: Dentley, David R.
APPLICANT: Dentley, David R.
APPLICANT: D'CONDOR, TIMOTHY
ITILE OF INVENTION: The Semaphorin Gene Family
NUMBER OF SEQUENCES: 100
CORRESPONDENCE ADDRESS:
ADDRESSEE: SCIENCE & TECHNOLOGY LAW GROUP
STREET: 268 Bush Street, Suite 3200
CITY: San Francisco
US-09-391-741A-9
US-09-620-312D-562
US-08-985-950-5
US-08-985-950-5
US-09-346-049-5
US-09-346-049-11
US-08-99-774A-11
US-08-99-774A-11
US-09-148-545-10
US-09-148-545-46
US-09-620-312D-7
US-09-620-312D-7
US-09-620-312D-7
US-09-621-763-7
US-09-621-763-7
US-09-621-763-7
US-09-621-763-7
US-09-221-235-1
US-09-221-235-1
US-09-221-235-1
US-09-221-235-1
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Pred. No. 0;
                                                                                                                                                                                                                                                                                                                                                                                               Sequence 53, Application US/08121713D
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SEQUENCE CHARACTERISTICS:
LENGTH: 2601 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
 11059
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), LOCATION: 16..2331
US-08-121-713D-53
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Best Local Similarity
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    Sequence 53, Appl
Sequence 53, Appl
Sequence 1, Appli
Sequence 5, Appli
Sequence 6, Appli
Sequence 7, Appli
Sequence 1, Appli
Sequence 4, Appli
Sequence 4, Appli
Sequence 23, Appli
Sequence 23, Appli
                                                                                                                         (without alignments)
7749.288 Million cell updates/sec
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                                                                                                                                                                                                   1 aatcttttatttatcgatg......aggcttttttcctaatacc 2709
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Sequence 1, A
Sequence 1, A
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Sequence 33,
Sequence 1,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /cgn2_6/ptodata/2/ina/5A_COMB.seq:*
/cgn2_6/ptodata/2/ina/5B_COMB.seq:*
/cgn2_6/ptodata/2/ina/6A_COMB.seq:*
/cgn2_6/ptodata/2/ina/FD_COMB.seq:*
/cgn2_6/ptodata/2/ina/PCTUS_COMB.seq:*
/cgn2_6/ptodata/2/ina/PCTUS_COMB.seq:*
                GenCore version 5.1.6
(c) 1993 - 2004 Compugen Ltd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-08-121-713D-53
US-08-835-268-53
US-08-9060-653
US-08-060-610-53
PCT-0894-10151A-53
US-08-136-10151A-53
US-08-136-10151A-53
US-08-136-10151A-53
US-08-136-10151A-53
US-08-136-10151A-53
US-08-136-10151A-53
US-09-139-1741A-23
US-09-391-741A-23
US-09-391-741A-25
US-09-391-741A-23
US-09-391-7445-1
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US-09-780-641-1
US-09-833-381-1211
                                                                                                                                                                                                                                                                                                                                            Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                              682709 seqs, 277475446 residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SUMMARIES
                                                                                                        September 23, 2004, 01:33:05
                                                                         - nucleic search, using sw model
                                                                                                                                                                                                                                                                                                                                                                                                                          Post-processing: Listing first 45 summaries
                                                                                                                                                                                                                                 OLIGO_NUC
Gapop_60.0 , Gapext 60.0
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Maximum DB seq length: 200000000
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2709
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Match Length
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26601
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Score

Result

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1273 CAGGGATGGACCCAACTATCAATGGGTGCCTTATCAAGGAGAGTCCCCTATCCACGGCC 1332	1333 AGGAACTIGICCCAGGAAAACAITIGGIGGIHITIGACICIACAAAGGACCITCCTGATGA 1392 	naccittgcaagaagtcatccagccatgtacaatccagtgtttcctatgaacaa 145 	3 TCGCCCAATAGTGATCAAAACGGATGTAAATTATCAATTTACACAAATTGTCGTAGACCG 151 	3 AGTGGATGCAGAAQATGGACAGTATGATGTTATGTTTATCGGAACAGATGTTGGGACCGT 157	3 TCTTAAAGTAGTTTCAATTCCTAAGGAGACTTGGTATGATGATTTAGAAGAGGTTCTGCTGGA 163	1633 AGAAATGACAGTTTTCGGGAACCGACTGCTATTTCAGCAATGGAGCTTTCCACTAAGCA 1692 	1693 GCAACAATTATATTGGTTCAACGGCTGGGGTTGCCCAGGTCCCTTTACACGGGTGTGA 1752 	53 T 69	4 4	1873 AAGAATGGAGACCACTGACTCACTGTTCAGACTTACACCATGATAATCACCATGGCCA 1932 	1933 CAGCCCTGAAGAGAGAATCATCTATGGTGTAGAATAGTAGCACATTTTGGAATGCAG 1992	1993 TCCGAAGTCGCAGAGAGCGCTGGTCTATTGGCAATTCCAGAGGCGAAATGAAGAGCGAAA 2052 	2053 AGAAGAGATCAGAGGGATGATCATATCATCAGGACAGATCAAGGCCTTCTGCTACGTAG 2112 1869 AGAAGAGATCAGAGTGATGATCATATCATCAGGACAGATCAAGGCCTTCTGCTACGTAG 1928	2113 TCTACAACAGAAGGATTCAGGCAATTACCTCTGCCATGCGGTGGAACATGCGTTCATACA 2172	2173 AACTCTTCTTAAGGTAACCCTGGAAGTCATTGACACAGAGCATTTGGAAGAACTTCTTCA 2232 [2233 TAAAGATGATGAGAGATGGCTCTAAGACCAAAGAAATGTCCAATAGCATGACACCTAG 2292	3 CCAGAAGGTCTGGTACAGAGACTTCATGCAGCTCATCAACCACCCCAATCTCAACGAT 235 	
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3; Conservative 0; Mismatches 0; Indels 0; Gaps CTGCAGCATGATAACTAAGAATTGATTATTCTGGGGAGTATTAACTAAGG 2	9 CIGCAGCATGGGCTGGTTAACTAGGATTGTCTGTGTGGGGAGGTATTACTTAC	A ANTOTTIGGAATCCAACAATGATGATCATTTCAATGGCTTGGCCAACAGCTCCAGTTATCA 37.	TACCTTCCTTTTGGATGAGATGAGTTGGTGTTGTTGAGCAAAGGATCACATATT	TTCATTCCACCTGGTTAATATCAAGGATTTTCAAAGGATTGTGGGCCAGTATCTTACAC 49:	CAGAAGAAGAAGAAGAAGAAGAAGAAGAAGAAGAAGAAGA	CAAGGTACTTAAGGCATATAATCAGACTTGTACGCTGTGGAACGGGGGTTTTCA 61.	CANGGIACTIAAGCCATATATCAGACTCACTTGIACGCCTGTGGGAACGGGGGGTTTTCA 42 TCCAATTTTGCACCTACATTGAAATTGGACATCATCCTGAGGACAATATTTTTAAGCTGGA 67 TCCAATTTTGAATTTGAAATTGGACATCATCCTGAGGACAATATTTTTAAGCTGGA 67 TCCAATTTGAATTTGAAATTGAACTATTCTAATTGAAATTGAAGATGAA	429 TCCAATTTGCACCTACATTGAAATTGGACATCCTGAGGACAATATTTTTAAGCTGGA 488 673 GAACTCACATTTTGAAAACGGCCGTGGGAAGTCCATATGACCCTAAGCTGCTGACAGC 732 1	GANCICACATTITAAAAAGGGCCGTGGGAAAGAGTCCATAATCCTAAGGTGCTGACGGC 54 ATCCCTTTTAATAGATGGAGAATTATACTCTGGAACTGCAGGCTGATTTTATGGGGCGAGA 79 ATCCCTTTTAATAGATGGAAATTATACTCTGGAACTGCAGCTGATTTTATGGGGCGAGA 6 ATCCCTTTTAATAGATGGAATTATAATCTCTGGAACTGCAGCTGAATTTATAGGGGCGAGA 6	TICCITITATION OF THE CONTROL OF THE	CAGGIGGICAL COMMITTE TO THE PROCESS OF THE PROCESS	CAGGIGGI CARIGATACTITITICTICGI GACCACCI CATCI CAGAGAGIGACARICC 72 TGAAGATGACAAAGTATACTITITICTICGI GAAAAAAGGAAAAAAAAAAAAAAAAAAAAAAAAAA	TORAGE: TORACTANGE TOTAL CONTINUATION OF AN INCOME OF A SARAGE TOTAL CONTINUATION OF A SARAGE TOTAL CONTINUATION OF A SARAGE CONTINUATION	######################################	AAATGGCATTGACACTCATTTGATGAACTGCAGATGTATTCCTAATGAACTTTAAAGA 11	AAATGGCATTGACACTCATTTTGATGAACTGCAGGATGTAFTCCTAATGAACTTTAAAGA 9 TCCTAAAAATCCAGTTGTATATGGAGTGTTTACGACTTCCAGTAACATTTTCAAGGGTC 1	AGGGATA ATGCCC	

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TCCTAAAAATCCAGTTGTATATGGAGTGTTTACGACTTCCAGTAACATTTTCAAGGGATC 1212
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                                                                                                                                                                                                                                                                                                  129 AATGTTGGAATCCAACAATGTGATCACTTTCAATGGCTTGGCCAACAGCTCCAGTTATCA 188
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Matches 2508; Conserv
   ; LOCATION:
US-08-835-268-53
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GGATGAGTTCTGTGAACAAGTTTGGAAAAGGGAACGAAAAAAAGGCCAGG
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CITY: USA
ZIP: QA
COUNTRY: USA
ZIP: 94104
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATCHTIN PC-DOS/MS-DOS
SOFTWARE: PATCHTIN PC-DOS/MS-DOS
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/835,268
FILING DATE:
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
APPLICANT: Goodman, Corey S.
APPLICANT: Kolodkin, Alex L.
APPLICANT: Matthes, David R.
APPLICANT: Dentley, David R.
APPLICANT: O'CONDOY. Timothy
TITLE OF INVENTION: The Semaphorin Gene Family
NUMBER OF SEQUENCES: 100
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APPLICATION NUMBER: US/08/121,713
FILING DATE: 13.55P-1993
ATTORNEY/AGENT INFORMATION:
NAME: Osman, Richard A.
REGISTATION NUMBER: 36,627
REFERENCE/DOCKET NUMBER: B94-002-1
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
ADDRESSEE: SCIENCE & TECHNOLOGY LAW
STREET: 268 Bush Street, Suite 3200
CITY: San Francisco
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US-08-835-268-53
; Sequence 53, Application US/08835268
; Patent No. 5807826
; Patent No. 5807826
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(415) 343-4342
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SEQUENCE CHARACTERISTICS:
LENGTH: 2601 base pairs
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MOLECULE TYPE: cDNA
FEATURE:
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1213 AGCGTGTGTATGTGTGTGTGTGTGTGTGTGTGTGTGTGTG	1613 AGANATGACNGTTTTTGGGAACCGACTGCTATTTCAGAATGAGCTTTCCACTAAGCA 1508 1619 GGAACACTATTTTGGGAACCGACTGCTATTTTCAGCAATGGAGCTTTCCACTAAGCA 1508 1619 GGAACACTATATTTGGGAACCGCTGGGATTGCCAATGGACTTTCCACTAAGCA 1508 1619 GGAACACTATATTTGGTTCAACGGCTGGGGTTGCCCAGCTCCCTTTAACACGGTGTGA 1568 1751 FATTACGGAAAGCGTTGTCAACTGGTTGCCTCGCCCAGACCCTTACTGTGGTA 1568 1811 GGTTCTGCATGTTCTGGTTATTTCCCACTCGCCCAGACCCTTACTGTGGTA 1688 1812 GGTTCTGCATGTTCTGGTTATTTCCCACTCGCCCAGACCCTTACTGTGGTATT 1972 1619 GGTTCTGCATGTTCTGGTTATTTCCCACTCGCCAGACCCAAACACACAC

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2109 CCAGAAGGTCTGGTACAGAGACTTCATGCAGCTCATCAACCACCCCAATCTCAACACGAT 2168
2293 CCAGAAGGTCTGGTACAGAGTTCATGCAGCTCATCAACCACCCCCAATCTCAACACGAT 2352
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EBDIUM TYPE: Floppy disk
DEPOTER: IBM PC compatible
PERATING SYSTEM: PC-DOS/MS-DOS
OFTWARE: Patentin Release #1.0, Version #1.25
PPLICATION DUMBER: US/09/060,692
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PLICANT: Kolodkin, Alex L.

PLICANT: Matthes, David

PLICANT: Bentley, David R.

PLICANT: O'Connor, Timothy

PLES OF INVENTION: The Semaphorin Gene Family

RERR OF SEQUENCES: 100

RESPONDENCE ADDRESS:
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TREET: 268 Bush Street, Suite 3200
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CLASSIFICATION: 514
CLASSIFICATION: 514
RIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/121,713
FILING DATE: 13-SEP-1993
FILING DATE: 11-SEP-1993
FILING DATE: 13-SEP-1993
FILING DATE
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ce 53, Application US/09060692
. No. 5935865
AL INFORMATION:
"LICANT: Goodman, Corey S.
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UENCE CHARACTERISTICS:
ENGTH: 2601 base pairs
YPE: nucleic acid
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IP: 94104
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1989 AACTCTTCTTAAGGTAACCCTGGAAGTCATTGACACAGAGCATTTGGAAGAACTTCTTCA 2048
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                                                 2233 TAAAGATGATGATGGAGATGCTCTAAGACCAAAGAAATGTCCAATAGCATGACACCTAG
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/833,391
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Goodman, Corey S.
APPLICANT: Kolodkin, Alex L.
APPLICANT: Matches, David
APPLICANT: Bentley, David R.
APPLICANT: O'Connor, Timothy
TITLE OF INVENTION: The Semaphorin Gene Family
NUMBER OF SEQUENCES: 100
CORRESPONDENCE ADDRESS:
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REGISTRATION NUMBER: 36,627
REFERENCE/DOCKET NUMBER: B94-002-1
TELECOMMUNICATION INFORMATION:
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/121,713
FILING DATE: 13.5EP-1993
ATTORNEY/AGENT INFORMATION:
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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Patent No. 6013781
GENERAL INFORMATION:
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy of
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100.0%; Pred. No. 0;
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TELEX:
INFORMATION FOR SEQ ID NO: 53
SEQUENCE CHARACTERISTICS:
LENGTH: 2601 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
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Best Local Similarity 100.(
Matches 2508; Conservative
                                                                                                                                                            16..2331
                                                                                            TOPOLOGY: linear
MOLECULE TYPE: cDNA
FEATURE:
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US-08-833-391-53
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1033 TCTGGTGAATAAATGGACAACATTCCTCAAAGCTCGTCTGATTTGCTCAGGTCC 1092 1093 AAATGGACATTAAATGAACATTCCTCAAAGCTCGTCTGATTTGCTCAGGTCC 008 1093 AAATGGCATTGAACACTTTTTGATTGATTCCTCAGTACTTTAAAGA 1152 909 AAATGGCATTGAACACTCATTTTGATGAACTTCAGGATTTCCTAAACATTTAAAGA 1152 1153 TCCTAAAAATCCAGTTGTATTGATGAACTTCAGGATGATTTCCAGTAACTTTTAAAGA 168 1153 TCCTAAAAATCCAGTTGTATTGATGAAGTGTTTTACGACTTTTCAAGGGATC 1028 1213 AGCCGTGTGTATTATAGCATGATTTTACGACTTCCAGTAACATTTTCAAGGGATC 1028 1213 AGCCGTGTGTATTATAGCATGATGATGAGAAGGGTTCCTTGGTCCATTTCAAGGGATC 1028 1213 AGCCGTGTGTATTATAGCATGATGATGAGAAGGGTTCCTTGGTCCATTTCAAGGGATC 1028 1214 AGCCGTGTGTATTATAGCATGATGATGAGAAGGGTCTTCCTTGGTCCATTTCCAGGGATC 1028 1315 AGCCGTGTGTATATAGCATGATGATGATGAGAAGGGTCCTTTCCTTGGTCCATTTCCAGGGATC 1028 1316 AGCCGTGTGTATTATAGCATGATGATGATGAAGGAACTCCTTGGTCCATTTCCAGGGATC 1028 1317 CCAGGATGAACATTTAAACGTTTTAAACATTTAAAGAAGACCCTTATCCAGGCC 1148 1318 AGCCAATTGTCCCAGCAAACATTTGATGAGAAGAACCCTTATCCATGAACAA 1268 1319 TCGTCCAATAGTGATCAAAACATTTTGATCAAAAGAACCCTTACTAGAACAA 1268 1329 AGTGGATGAACAAAACATTTAGATGATAAAACAAAACAA	1633 AGAANTGACAGTTTTTCGGGAACCCACTGCTATTTCAGCAATGGAGCTTTCACCTAAGCA 1508 1449

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973 AAAAGCTACTCACGCTAGAATAGGTCAGATATGCAAGAATGACTTTGGAGGCACAGAAG 1032
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REGISTRATION NUMBER: 36,627
REPERENCE/DOCKET NUMBER: B94
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 343-4341
TELEFAX: (415) 343-4342
                                                                                                                INFORMATION FOR SEQ ID NO: 53:
SEQUENCE CHARACTERISTICS:
LENGTH: 2601 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                 ) NAME/KEY: CDS
; LOCATION: 16..2331
US-09-060-610-53
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92.6%; Score 2508; C
Best Local Similarity 100.0%; Pred. No. 0;
Matches 2508; Conservative 0; Mismatches
TELEPHONE: (415) 781-1989
TELERA: (415) 398-3249
TELEA: 910 277299 FHT UR
INFORMATION FOR SEQ 1D NO: 53:
SEQUENCE CHARACTERISTICS:
LENGTH: 2610 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
TOPOLGY: linear
MOLECULE TYPE: CDNA
FEATURE:
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                                                                                                                                                                                                                              ; NAME/KEY: CDS
; LOCATION: 16..
PCT-US94-10151A-53
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                                                                                                              1929 TCTACAACAGAAGGATTCAGGCAATTACCTCTGCCATGCGGTGGAACATGGGTTCATACA
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                                    2173 AACTCTTCTTAAGGTAACCCTGGAAGTCATTGACACAGAGCATTTGGAAGAACTTCTTCA
                                                                                                                                                                                                2233 TAAAGATGATGGAGATGGCTCTAAGACCAAAGAAATGTCCAATAGCATGACACCTAG
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                                                                                    TCTACAACAGAAGGATTCAGGCAATTACCTCTGCCATGCGGTGGAACATGGGTTCATACA
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GENERAL INFORMATION: The Regents of the University of California APPLICANT: The Regents of the University of California TITLE OF INVENTION: The Semaphorin Gene Family NUMBER OF SEQUENCES: 66
CORRESPONDENCE ADDRESS:
ADDRESSEE: FLEHR HOHBACH TEST ALBRITTON & HERBERT CITY: San Francisco Center, Suite 3400
STREET: 4 Embarcadero Center, Suite 3400
STREET: 5AP FRANCISCO COMPATEILE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER READBLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: 1BM PC Compatible
OPERATING SYSTEM: PCT/US94/10151A
FILING DATE: 13-SEP-1994
CLASSIFICATION:
NAME: OSMAN: Richard A.
REFERENCE/DOCKET NUMBER: FP-58750-PC/RAO
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
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1078 CCATGGCCAGGCCCTGAAGAGAAATCATCTATGGTGTAGAAATAGTAGCACATTTTT 1137
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                                                                                                                                                           ACAAGATATAAGAAATGGAGACCCACTGACTCACTGTTCAGACTTACACCATGATAATCA 1923
                                                                                                                                                                                                                                                                                                                                                                                                                                        1138 ggaangcagnccgaagncgcagagagcgcrcgrcrarrcgcaarrccagaggcgaaarga 1197
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AGAGCGAAAAGAAGAGATCAGAGGATGATCATATCATCAGGACAGATCAAGGCCTTCT 1257
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                                                                                                                               1804 TGCTTGGGATGGTTCTGCATGTTCTCGCTATTTTCCCACTGCAAAGAGACGCACAAGACG
                                                                                                                                                                                                                                                                                                             CCATGGCCACAGCCCTGAAGAGAGAAATCATCTATGGTGTAGAGAATAGTAGCACATTTTT
CACTAAGCAGCAACAACTATATATTGGTTCAACGGCTGGGGGTTGCCCAGCTCCCTTTACA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 5, Application US/08786531B
Sequence 5, Application US/08786531B
Patent No. 6541197
GENERAL INFORMATION:
APPLICANT: Levy, John P.
APPLICANT: Levy, John P.
APPLICANT: Rang, Suming
APPLICANT: Seregime 1
TITLE OF INVENTION: Vehicles for Stable Transfer of Green
TITLE OF INVENTION: Phorescent Protein Gene and Methods of Use for SNUMBER OF SEQUENCES: 10
CORRESPONDENCE ADDRESS: 10
CORRESPONDENCE ADDRESS: 10
CORRESPONDENCE ADDRESS: 2arley, McKee, Thomte, Voorhees & Sease
                                                                                        898 ¢¢¢¢t¢t¢atatttac¢¢aaaa¢¢¢t¢t¢ct¢a¢tett¢cct¢¢cccqaqaccttact¢
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPATING SYSTEM: PC-DCS/MS-DCS
SOFTWARE: RatentIn Release #1.0, Version #1.30
CURRINT APPLICATION DATA:
FILING DATE: 21-0AN-1997
CLASSIFICATION: DATA:
FILING DATE: 21-0AN-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/010371
FILING DATE: 22-JAN-1996
ATTORNEY/AGENT INFORMATION:
NAME: NOBEL, Heldi S.
REGISTRATION NUMBER: 37,719
REFERENCE/DOCKET NUMBER: Hgtri
TELEPHONE: 515-288-3667
TELEPHONE: 515-288-138
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE: Zarley, McKee, The STREET: 801 Grand Suite 3200 CIIY: Des Moines STATE: Iowa COUNTRY: United States
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-08-786-531B-5/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GCCAGGTCCAAATGGCATTGACACTCATTTTGATGAACTGCAGGATGTATTCCTAATGAA 1143
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ACACTCTGGAAAAGCTACTCACGCTAGAATAGGTCAGATATGCAAGAATGACTTTGGAGG 1023
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 58 IGACAATCCIGAAGAIGACAAAGIATACITITICITICGIGAAAAIGCAATAGAIGGAGA 117
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       118 ACACTCTGGAAAAGCTACTCACGCTAGAATAGGTCAGATATGCAAGAATGACTTTGGAGG 177
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                                                                                                                                                                                                                                                                                                                                       Length 1481
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                                                                                                                                                                                                                                                                                                                                       DB 1;
                                                                                                                                                                                                                                                                                                                                  Query Match
46.3%; Score 1253; Di
Best Local Similarity 99.9%; Pred. No. 0;
Matches 1303; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                            904 TGACAATCCTGAAGATGACAAAGTATACT
    TELECOMMUNICATION INFORMATION:
TELEPHONE: 215-568-3100
TELEPRAX: 215-568-3439
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
                                                                                                             LENGTH: 1481 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
TOPOLOGY: both
MOLECULE TYPE: cDNA
                                                                                                                                                                                                                                              CDS
50..1480
                                                                                                                                                                                                                                              NAME/KEY:
LOCATION:
                                                                                                                                                                                                                                                                          ;
US-08-136-922-1
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC comparatible
COMPUTER: TOP C comparatible
COMPUTER: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/786,531B
FILING DATE: 21-JAN-1997
CLASSIFICATION:
                                                                4254 AATCITITATITATCGATGITAACAAGCIT 4224
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            4330 AArchrharhrharccarch 4309
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APPLICATION NUMBER: US 60/010371
FILING DATE: 22-JAN-1996
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 AATCTTTTATTTATCGATGTT 22
                                                                                                                                                                                                                           Sequence 3, Application US/09645004; Patent No. 6608238
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; OTHER INFORMATION: Plasmid pLNMX US-09-645-004-3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Nebel, Heidi S.
REGIESTRATION NUMBER: 37,715
REFERENCE/DOCKET NUMBER: bg
TELECOMMUNICATION INFORMATION.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CITY: Des.
STATE: Iowa
COUNTRY: United States
71P: 50309
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: DNA ORGANISM: Artificial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-08-786-531B-4/c
                                                                                                                                                                                                US-09-645-004-3/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FEATURE:
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US-08-786-531B-6/C

Sequence 6, Application US/08786531B

Fetent No. 654197

GENERAL INFORMATION:

APPLICANT: Link, Charles J.

APPLICANT: Link, Charles J.

APPLICANT: Link, Charles J.

APPLICANT: Seregian Tation

TITLE OF INVENTION: Vehicles for Stable Transfer of Green

TITLE OF INVENTION: Pluorescent Protein Gene and Methods of Use for SEQUENCES: 10

CORRESPONDENCES: 10

CORRESPONDENCES: 2016-8.

ADDRESSEE Zatles

STREET: BOIL Grad States

STREET: BOIL Grad States

CONTY: Des Moines

CONTY: Des Moines

STREET: BOIL Grad States

CONTY: IBM PC compatible

CONTY: IBM PC compatible

CONTY: REPEADABLE FORM:

MEDIOM TYPE: Ploppy disk

CONTY: IBM PC compatible

CONTY: STREET: 40:000

CONTY: 1000

CONTY: 10
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100.0%; Pred. No. 4.3e-05;
ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                           Length 7160;
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                                                                                                                                                                                                                                                                                                                                                              Query Match
1.1%; Score 31; DB 4; L.
Best Local Similarity 100.0%; Pred. No. 4.3e-05;
Matches 31; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          4179 AATCITITATITITATCGATGTTAACAAGCIT 4149
| INFORMATION FOR SEQ ID NO: 5:
| SEQUENCE CHARACTERISTICS:
| LENGTH: 7160 base pairs
| TYPE: nucleic acid
| STRANDEDNESS: single
| TOPOLOGY: linear
| MOLECULE TYPE: other nucleic acid
| HYPOTHETICAL: NO
| ANTI-SENSE: NO
| JUS-08-786-531B-5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      other nucleic acid
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TELEPAX: 515-288-1338
INFORMATION FOR SEQ ID NO: ~6:
SEQUENCE CHARACTERISTICS:
LENGTH: 7235 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity 100.
Matches 31; Conservative
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STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           linear
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APPLICANT: Gavora, Jan S.
APPLICANT: Falconer, Marcia M.
APPLICANT: Palconer, Marcia M.
APPLICANT: Nguyen, Thuy H.
APPLICANT: Benkel, Bernhard F.
TITLE OF INVENTION: Trans-Somatics With Gene Transfer Into Mammary Epithelial Cells FILE REFERENCE: GALA-06402.
CURRENT APPLICATION NUMBER: US/09/645,004
CURRENT FILING DATE: 2000-08-24
NUMBER OF SEQ ID NOS: 4
SOFTWARE: PatentIn version 3.2
SEQ ID NO 3
LIBRORDH: 7311
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Sequence 4, Application US/08786531B
Patent No. 654197
GENERAL INFORMATION:
APPLICANT: Link, Charles J.
APPLICANT: Link, Charles J.
APPLICANT: Seregina, Tatiana
TITLE OF INVENTION: Vehicles for Stable Transfer of Green
TITLE OF INVENTION: Vehicles for Stable Transfer of Green
TITLE OF INVENTION: Vehicles for Stable Transfer of Green
TITLE OF INVENTION: Vehicles for Stable Transfer of Green
TITLE OF INVENTION: Vehicles for Stable Transfer of Green
TITLE OF INVENTION: Vehicles for Same
NUMBER OF SEQUENCES: 10
CORRESPONDENCE ADDRESS:
ADDRESSEE: Zarley, McKee, Thomte, Voorhees & Sease
STREET: 801 Grand Suite 3200
CITY: Des Moines
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0; Gaps
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Indels
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    Mismatches
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, OTHER INFORMATION: Plasmid pLNMi2X
US-09-645-004-4
                                                                                                                                                                                     Sequence 4, Application US/09645004
Patent No. 6608238
GENERAL INFORMATION:
    22; Conservative
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; LOCATION: (172)...(807)
US-09-391-741A-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: DNA ORGANISM: Artificial
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ORGANISM: Zea mays
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US-09-645-004-4/c
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    Matches
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US-08-786-531B-1/C
US-08-786-531B-1/C
Sequence 1, Application US/08786531B
Fatent No. 6541197
GENERAL INFORMATION:
APPLICANT: Link, Charles J.
APPLICANT: Levy, John P.
APPLICANT: Seregina, Tatiana
TITLE OF INVENTION: Vehicles for Stable Transfer of Green
TITLE OF INVENTION: Pluorescent Protein Gene and Methods of Use for Same
NUMBER OF SEQUENCES: 10
CORRESPONDENCE ADDRESS:
ADDRESSEE: Zarley, McKee, Thomte, Voorhees & Sease
STREET: 801 Grand Suite 3200
CITY: Des Moines
STARE: Inwa
TITLE OF INVENTION: Pluorescent Protein Gene and Methods of Use for Same
NUMBER OF SEQUENCES:
ADDRESSEE: Aarley, McKee, Thomte, Voorhees & Sease
STREET: Bol Grand Suite 3200
CITY: Des Moines
STARE: Inwa
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                                                                                                                                                                                                                                                                             0.8%; Score 22; DB 4; Length 7352; 00.0%; Pred. No. 0.97;
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                                                                                                                                                                                                                                                                                                                       0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM FC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/786,531B
FILING DATE: 21-JAN-1997
CLASSIPICATION: A35
PRIOR APPLICATION NUMBER: US 60/010371
FILING DATE: Z2-JAN-1996
ATORNEY/AGENT INFORMATION:
NAME: NAME: NAME: NAME: 37,719
REGISTRATION NUMBER: 37,719
                                                                                                                                                                                                                                                                                                 Pred. No. 0.9
Mismatches
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100.0%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                        4371 AArcritaririnarcearcri 4350
                                                                                                                                             TOPOLOGY: linear MOLECULE TYPE: other nucleic acid HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    other nucleic acid
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Conservative 0;
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 515-288-3667
TELEPAX: 515-288-1338
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
TELEPHONE: 515-288-3667
TELEFAX: 515-288-138
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
LENGTH: 7352 base pairs
LYPE: nucleic acid
STRANDEDNES; single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 7353 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
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Matches 22; Conserv
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Best Local Similarity
                                                                                                                                                                                                                ; ANTI-SENSE: NO
US-08-786-531B-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MOLECULE TYPE:
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APPLICANT: Gavora, Jan S.
APPLICANT: Gavora, Jan S.
APPLICANT: Gavora, Jan S.
APPLICANT: Marcia M.
APPLICANT: Mayen, Thuy H.
APPLICANT: Benkel, Bernhard F.
TITLE OF INVENTION: Trans-Somatics With Gene-Transfer Into Mammary Epithelial Cells
FILE REPERBRYEE: GALA-06402
CURRENT APPLICATION NUMBER: US/09/645,004
CURRENT FILING DATE: 2000-08-24
NUMBER OF SEQ ID NOS: 4
SEQ ID NO 4
LENGTH: 7885
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 23, Application US/09391741A
Patent No. 655532
GENERAL INFORMATION:
APPLICANT: DUVICK, Jonathan P.
APPLICANT: Sharma, Yogesh Kumar
TITLE OF INVENTION: Rac-Like Genes and Methods of Use
FILE REFERENCE: 0866D;
CURRENT FILING DATE: 1999-09-08
PRIOR PLICATION NUMBER: 60/111,919
PRIOR FILING DATE: 1998-12-11
PRIOR FILING DATE: 1998-12-11
PRIOR FILING DATE: 1998-09-14
PRIOR FILING DATE: 1999-09-08
NUMBER OF SEQ ID NOS: 51
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 23
LENGTH: 1062
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3 GTCGACCCACGCGTCCGGGAG 23

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Query Match

0.8%; Score 21; DB 4; Length 1062;
Best Local Similarity 100.0%; Pred. No. 3;
Matches 21; Conservative 0; Mismatches 0; Indels
RESULT 15
US-09-391-741A-13
is Sequence 33, Application US/09391741A
is Patent No. 6555732
is GENERAL INFORMATION:
APPLICANT: Durvick, Jonathan P.
APPLICANT: Durvick, Jonathan P.
APPLICANT: Brana, Yogesh Kumar
ITILE OF INVENTION: Rac-Like Genes and Methods of Use
FILE REPERENCE: 0866D
CURRENT APPLICATION NUMBER: US/09/391,741A
CURRENT FILING DATE: 1999-00-08
is PRIOR FILING DATE: 1998-11-11
PRIOR PILING DATE: 1998-14
is PRIOR APPLICATION NUMBER: 09/391,741
is PRIOR APPLICATION NUMBER: 09/391,741
is PRIOR FILING DATE: 1999-09-08
is NUMBER OF SEQ ID NOS: 51
is SOFTWARE: FeatSEQ for Mindows Version 3.0
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Search completed: September 23, 2004, 05:10:25 Job time : 204 secs

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APPLICANT: Karen GLATT
TITLE OF INVENTION: Nucleic Acid Molecules and Proteins For The Identification,
TITLE OF INVENTION: Assessment, Prevention, and Therapy of Ovarian Cancer
FILE REFERENCE: MRI-030
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   284, App
10, Appl
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   seduence Sed
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US-10-097-340-284
US-10-262-538-10
US-10-267-632-64
US-10-36-638-10
US-10-36-6318-10
US-10-247-671-164
US-10-29-946-374-310
US-10-20-915-348
US-10-20-915-348
US-10-20-915-348
US-10-20-915-348
US-10-20-916-348
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US-10-10-20-914-348
US-10-10-176-749-348
US-10-176-749-348
US-10-176-749-348
US-10-180-550-348
US-10-180-550-348
US-10-180-550-348
US-10-180-550-348
US-10-180-550-348
US-10-194-460-348
US-10-194-460-348
US-10-194-460-348
US-10-194-460-348
US-10-196-755-348
US-10-199-304-348
US-10-199-304-348
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US-10-202-412-348
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Publication No. US20030087250A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: John MONAHAN
APPLICANT: John MONAHAN
APPLICANT: Sebastian HOERSCH
APPLICANT: State of KOVATS
APPLICANT: State of KOVATS
APPLICANT: Rechel E MEYERS
APPLICANT: Rechel E MEYERS
APPLICANT: Peter OLANDT
APPLICANT: Peter OLANDT
APPLICANT: APPLICANT: Gordon B. MILLS
APPLICANT: Gordon B. MILLS
APPLICANT: Robert C. BAST, Jr.
APPLICANT: Robert C. BAST, Jr.
APPLICANT: Robert C. BAST, Jr.
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Xumei ZHAO
Karen GLATT
      US-10-097-340-284
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   3; Search time 303 Seconds (without alignments) 5742.244 Million cell updates/sec
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| cgn2_6/ptodata/2/pubpaa/US07_PUBCOMB.pep:*
2: /cgn2_6/ptodata/2/pubpaa/US06_NEW PUB.pep:*
3: /cgn2_6/ptodata/2/pubpaa/US06_NEW PUB.pep:*
4: /cgn2_6/ptodata/2/pubpaa/US06_PUBCOMB.pep:*
5: /cgn2_6/ptodata/2/pubpaa/US06_PUBCOMB.pep:*
6: /cgn2_6/ptodata/2/pubpaa/US08_NEW PUB.pep:*
7: /cgn2_6/ptodata/2/pubpaa/US08_NEW PUB.pep:*
8: /cgn2_6/ptodata/2/pubpaa/US08_NEW PUB.pep:*
9: /cgn2_6/ptodata/2/pubpaa/US08_PUBCOMB.pep:*
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11: /cgn2_6/ptodata/2/pubpaa/US09_PUBCOMB.pep:*
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14: /cgn2_6/ptodata/2/pubpaa/US08_PUBCOMB.pep:*
15: /cgn2_6/ptodata/2/pubpaa/US08_PUBCOMB.pep:*
16: /cgn2_6/ptodata/2/pubpaa/US10_NEW PUB.pep:*
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                                      GenCore version 5.1.6
(c) 1993 - 2004 Compugen Ltd.
                                                                                                                                          protein search, using frame_plus_n2p model
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Maximum Match 100%
Listing first 45 summaries
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Xgapop 10.0 , Xgapext
Ygapop 10.0 , Ygapext
Fgapop 6.0 , Fgapext
Delop 6.0 , Delext
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Matches:
Conservative:
Mismatches:
Indels:
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CURRENT APPLICATION NUMBER: US/10/097,340
CURRENT FILING DATE: 2002-03-14
PRIOR PILING DATE: 2001-03-14
PRIOR FILING DATE: 2001-03-14
PRIOR FILING DATE: 2001-03-15
PRIOR FILING DATE: 2001-03-26
PRIOR APPLICATION NUMBER: 60/325,149
PRIOR FILING DATE: 2001-03-14
PRIOR FILING DATE: 2001-03-14
PRIOR FILING DATE: 2001-03-14
PRIOR FILING DATE: 2001-03-16
PRIOR FILING DATE: 2001-09-26
PRIOR FILING DATE: 2001-09-19
PRIOR FILING DATE: 2001-09-19
NUMBER OF SEQ ID NOS: 363
SOFTWARE: FASTSEQ fOR WINDOWS VERSION 4.0
SOFTWARE: FASTSEQ fOR WINDOWS VERSION 4.0
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100.00%
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85.80%
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CORGANISM: Homo sapiens
US-10-097-340-284
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Best Local Similarity:
Query Match:
DB:
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INFORMATION FOR SEQ ID NO: 54:
SEQUENCE CHARACTERISTICS:
LENGTH: 771 amino acid
TYPE: amino acid
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SEQUENCE DESCRIPTION: SEQ
US-10-067-632-54
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GlyProAsnTyrGlnTrpValProTyrGlnGlyArgValProTyrProArgProGlyThr 380
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
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APPLICATION NUMBER: US/10/067,632
FILING DATE: 04-Feb-2002
CLASSIFICATION: «Unknown»
PRIOR APPLICATION DATA:
FILING DATE: «Unknown»
FILING DATE: «Unknown»
APPLICATION NUMBER: 08/835,268
FILING DATE: «Unknown»
APPLICATION NUMBER: OB/835,268
FILING DATE: «Unknown»
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: Osman, Richard A.
REGISTRATION NUMBER: 36,627
REPERENCE/POCKET NUMBER: B94-002-1
TELEPHONE: (415)343-4341
                                                            761 ThrHisGluPheGluArgAlaProArgSerVal
                                                2480 ACCCACGAATTTGAGAGGCACCCAGGAGTGTC
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US-10-247-671-164
                                                                                                                                      CELLS
                                                           NS-10-147-671-164

1 Sequence 164, Application US/10247671

2 FUBLICATION NO. USZ0030194721A1

3 GENERAL INPORMATION:

APPLICANT: Mikita, Thomas

APPLICANT: Machew R.

TITLE OF INVENTION: GENES EXPRESSED IN TREATED FOAM CELL

APPLICANT: Raser, Matthew R.

TITLE OF INVENTION OF USCOLOGO US

CURRENT APPLICATION NUMBER: US/10/247,671

CURRENT APPLICATION NUMBER: 00/3-19

PRIOR FILING DATE: 2001-09-19

NUMBER OF SEQ ID NOS: 186

SOFTWARE: PERL PROGRAM

LENGTH: 71

TYPE: PRT

CURRENT APPLICATION OF USCOLOGO

SOFTWARE: PERL PROGRAM

SEQ ID NO 164

CORGANISM: Homo sapiens
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Matches:
Conservative:
Mismatches:
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                    201 IlePheArgThrLeuGlyHisHisHisProlleArgThrGluGlnHisAspSerArgTrp
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 CATITIGAAAACGGCCGTGGGAAGAGTCCATATGACCCTAAAGCTGCTGACAGCATCCCTT
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SM: Homo sapiens 769-3 Scores: 0	0-0 0-0 4-0 4-d H-	rGlyA PTTTA PPhel Phel ScrGC 	CTCAATGATCGAAGTTCATTAGTGCCCCCCCTCATCTCAGAGAGTGACAATCCTGAAGAT LEUASHASPPLOLYSPHEIJESETALAHISLEUIILESETGIUSETASPASHPYOGIUASP GACAAAGTATCTTTTCTCCGTGAAAATGCAATAGATGGAGAACCTCTGGAAAGCT ASPLYSVAITYTPHEPHEPHEAYGGIUASNAIAIILEASPGIYGIUHISSETGIYLYSALA ACTCACGCTAGAATAGGTCAGATGAAAAGAATAGATAGAGAAGAAGATCTGGAAAGCT
) ORGANI US-10-320- Alignment Pred. No.: Score: Percent Si Best Local Query Matc DB: US-09-774- Qy Db	6 8 6 8 6 8 6 6 6	4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4	 66666666666666666666666666666666666
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US-08-946-374-310 Sequence 310, Application US/09946374 PUBLICARTION NO. US20030073129A1 GREEAL INFORMATION: APPLICANT: Baker, Kevin P. APPLICANT: Bater, Kevin P. APPLICANT: Bater, Mevin P. APPLICANT: Baten, Dan L. APPLICANT: Fong, Sherman APPLICANT: Gadowski, Napoleone APPLICANT: Godowski, Paul J. APPLICANT: Godowski, Paul J. APPLICANT: Godowski, Paul J. APPLICANT: Godowski, Paul J.	Gurney, P Hillan, P Pan, Jame Pan, Jame Roy, Marc Smith, Vi Stewart, Tumas, De Watanabe, Williams, Wood,	FITLE REFERENCE: P2330PLCI CURRENT APPLICATION NUMBER: US/09/946,3 CURRENT FILING DATE: 2001-09-04 FRIOM APPLICATION NUMBER: 60/098716 FRIOM FILING DATE: 1998-09-01 FRIOM FILING DATE: 1998-09-01 FRIOM FILING DATE: 1998-09-01 FRIOM APPLICATION NUMBER: 60/098723 FRIOM FILING DATE: 1998-09-01 FRIOM FILING DATE: 1998-09-01	PRIOR FILING DATE: 1998-09-01 PRIOR FILING DATE: 1998-09-01 PRIOR PLING DATE: 1998-09-02	PRIOR PEPLICATION NUMBER: 00/095996 PRIOR APPLICATION NUMBER: 00/095998 PRIOR APPLICATION NUMBER: 00/095602 PRIOR APPLICATION NUMBER: 00/09602 PRIOR PILING DATE: 1998-09-09 PRIOR PILING DATE: 1998-09-09 PRIOR PILING DATE: 1998-09-09 PRIOR PILING DATE: 1998-09-09 PRIOR PILING DATE: 1998-09-10	FRIOR FILING DATE: 1936-09-10 FRIOR FILING DATE: 1986-09-10 FRIOR FILING DATE: 1986-09-10 FRIOR FILING DATE: 1996-09-10 FRIOR PRIOR DATE: 1996-09-10 FRIOR FILING DATE: 1996-09-10 FRIOR FILING DATE: 1996-09-10 FRIOR APPLICATION NUMBER: 60/099815 FRIOR APPLICATION NUMBER: 60/099815 FRIOR FILING DATE: 1996-09-10 FRIOR FILING DATE: 1996-09-10 FRIOR FILING DATE: 1996-09-10 FRIOR FILING DATE: 1998-09-10
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APPLICANT: Month, Mustandbe, Colin K.
APPLICANT: Month, Mulliam I.
APPLICANT: Zhang, Zemin K.
APPLICANT: Zhang, Zemin A.
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PRICA APPLICATION NUMBER: U0/10/206, 915
PRIOR APPLICATION NUMBER: U0/05266
PRIOR FILING DATE: 1997-09-18
PRIOR FILING DATE: 1997-10-24
PRIOR APPLICATION NUMBER: 60/063120
PRIOR FILING DATE: 1997-10-24
PRIOR APPLICATION NUMBER: 60/063120
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Godowski, Paul J.
Gurney, Austin L.
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 1580 GTAGTTTCAATTCCTAAGGAGACTTGGTATGATTTAGAAGAGGTTCTGCTGGAAGAAATG 1639
                                                                                                                                                                                                     GGGAAAGCGTGTGCTGAGTGTTGCCTCGCCCGAGACCCTTACTGTGCTTGGGATGGTTCT 1819
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                                                                                                                                  CTATATATTGGTTCAACGCCTGGGGTTGCCCAGCTCCCTTTACACCGGGTGTGATATTAC 1759
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                  538-GlyLysAlaCysAlaAspCysCysLeuAlaArgAspProTyrCysAlaTrpAspGlyAsn 557
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Sequence 348, Application US/10199670 Publication No. US20040033560A1 GENERAL INFORMATION:

US-10-199-670-348

Desnoyers, Luc Goddard, Audrey

APPLICANT: Baker, Kevin P.
APPLICANT: Chen, Jian
APPLICANT: Desnoyers, Luc
APPLICANT: Goddard, Audre)

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                                                                                                     572 AATCAGACICACTIGIACGCCTGIGGAACGGGGCTTITCAICCAATTTGCACCTACATT
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PRIOR FILING DATE: 2002-01-15
PRIOR APPLICATION NUMBER: 60/05926
PRIOR APPLICATION NUMBER: 60/05926
PRIOR APPLICATION NUMBER: 60/06226
PRIOR APPLICATION NUMBER: 60/06226
PRIOR PLING DATE: 1997-09-18
PRIOR APPLICATION NUMBER: 60/063120
PRIOR PLING DATE: 1997-10-17
PRIOR APPLICATION NUMBER: 60/063121
PRIOR APPLICATION NUMBER: 60/063121
PRIOR APPLICATION NUMBER: 60/063540
PRIOR FILING DATE: 1997-10-21
PRIOR PRIOR DATE: 1997-10-21
PRIOR APPLICATION NUMBER: 60/063540
PRIOR FILING DATE: 1997-10-28
PRIOR PLING DATE: 1997-10-28
PRIOR PLING DATE: 1997-10-28
PRIOR PLING DATE: 1997-10-28
PRIOR PLING DATE: 1997-10-28
PRIOR APPLICATION NUMBER: 60/063544
PRIOR APPLICATION NUMBER: 60/063544
PRIOR PLING DATE: 1997-10-28
PRIOR APPLICATION NUMBER: 60/063544
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PRIOR APPLICATION NUMBER: 60/063544
PRIOR APPLICATION NUMBER: 60/063544
PRIOR PLING DATE: 1997-10-28
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PRIOR APPLICATION NUMBER: 60/063544
PRIOR PLING DATE: 1997-10-28
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PRIOR APPLICATION NUMBER: 10/052586
Pan, James
Smith, Victoria
Watanabe, Colin K.
Wood, William I.
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Publication No. US20040038337A1
GENERAL INFORMATION:
APPLICANT: Baker, Kevin P.
APPLICANT: Chen, Jian
APPLICANT: Desnoyers, Luc
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Smith, Victoria
Watanabe, Colin K.
Wood, William I.
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Godowski, Paul J.
Gurney, Austin L.
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|LeuGlyProThrHisAspHisHisTyrIleArgThrAspIleSerGluHisTyrTrpLeu 238
                                                                                                                                                               AATGATCCAAAGTTCATTAGTGCCCACCTCATCTCAGAGAGTGACAATCCTGAAGATGAC 922
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      GAAATTGGACATCATCCTGAGGACAATATTTTTAAGCTGGAGAACTCACATTTTGAAAAC
                                            GGCCGTGGGAAGAGTCCATATGACCCTAAGCTGCTGACAGCATCCCTTTTAATAGATGGA
                                                        GAATTATACTCTGGAACTGCAGCTGATTTTATGGGGCGAGACTTTGCTATCTTCCGAACT
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LeuArgTyrLysAspTyrIleGlnIleLeuSerSerProAsnPhe---SerLeuAspGln 729
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GGAGACCCACTGACTCACTGTTCAGACTTACACCATGATAATCACCATGGCCACAGGCCCT
                                                                                                                                                                                                                                                                          GAAGAGAGAATCATCTATGGTGTAGAGAATAGTAGCACATTTTTGGAATGCAGTCCGAAG
                                                                                                                                                                                                                                                                                                748 ------ProlysTrpLysHisMetGlnGluMetLysLysLysArgAsnArgArg
                                                 GGGAAAGCGTGTGCTGGCTCGCCCGAGACCCTTACTGTGCTTGGGATGGTTCT
                                                                                                                         GCATGTTCTCGCTATTTTCCCACTGCAAAGAGGACGCACAAGACGACAAGAAAT
                                                                                                                                                                                                                                                                                                                                                  TCGCAGAGAGCGCTGGTCTATTGGCAATTCCAGAGGCGAAATGAAGAGCGAAAAGAGAG
                                                                                                                                                                                                                                                                                                                                                                             2060 ATCAGAGTGGATGATCATATCATCAGGACAGATCAAGGCCTTCTGCTACGAGACTACAA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUCLEIC
                                                                                                                                                                                                                      APPLICANT: WOOD, William I.
APPLICANT: Zhang, Zemin
TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2480 ACCCAC---GAATTTGAGAGGCCACCCAGGAGTGTC
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us-09-774-490-1.rapb

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99 AsnLysAsnPheLysLysIleTyrTrpProAlaAlaLysGluArgValGluLeuCysLys 118
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FILE REPERENCE: P3430R1C464
CURRENT APPLICATION: ACIDS ENCODING THE SAME
FILE REPERENCE: P3430R1C464
CURRENT APPLICATION NUMBER: US/10/201,858
PRIOR APPLICATION NUMBER: 10/052586
PRIOR FILING DATE: 1997-09-18
PRIOR FILING DATE: 1997-09-18
PRIOR PLING DATE: 1997-09-18
PRIOR APPLICATION NUMBER: 60/05266
PRIOR FLING DATE: 1997-09-18
PRIOR PILING DATE: 1997-10-18
PRIOR PILING DATE: 1997-10-24
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PRIOR FILING DATE: 1997-10-24
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Mismatches:
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US-10-201-858-348
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Best Local Similarity:
Query Match:
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1759 CTATATATTGGTTCAACGGCTGGGGGTTGCCCAGCTCCCTTTACACGGGTGTGATATTTAC 1759
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                                                                                                                                                                                                                                                      983 CACGCTAGAATAGGTCAGATATGCAAGAATGACTTTGGAGGGCACAGAAGTCTGGTGAAT 1042
                                                                                                                                                                                                                                                                                                                                                                                   103 GACACTCATTTTGATGAACTGCAGGATGTATTCCTAATGAACTTTAAAGATCCTAAAAAT 1162
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                                                                                                                                                                                                                                                                                                                    1043 AAATGGACAACATTCCTCAAAGCTCGTCTGATTTGCTCAGTGCCAGGTCCAAATGGCATT 1102
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1223 AIGTATAGCATGAGTGATGTGAAAGGTGTTCCTTGGTCCATATGCCCACAGGGATGGA 1282
                 379 AlaAspHisArgTrpValGlnTyrAspGlyArgIleProTyrProArgProGlyThrCys 398
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752 GAATTATACTCTGGAACTGCAGCTGATTTTATGGGGCGAGACTTTGCTATCTTCCGAACT 811
                                                                                                                                                                                        923 AAAGTATACTTTTTTTTCTTCCGTGAAAATGCAATAGATGGAGAACACTCTGGAAAAGCTACT 982
                                                                                  | 1283 CCCAACTATCAATGGGTGCCTTATCAAGGAAGGAGGGCCCTATCCACGGCCAGGAACTTGT
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                                                                                                                            863 AATGATCCAAAGTTCATTAGTGCCCACCTCATCTCAGAGAGTGACAATCCTGAAGATGAC
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                                                               812 CTTGGG------CACCACCACCCAATCAGGACAGAGCAGCATGATTCCAGGTGGCTC
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119 LeuAlaGlyLysAspAlaAsnThrGluCysAlaAsnPheIleArgValLeuGlnProTyr 138
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PRIOR APPLICATION NUMBER: 60/059266

PRIOR FILING DATE: 1997-09-18

PRIOR PRING DATE: 1997-09-18

PRIOR PILING DATE: 1997-10-17

PRIOR PILING DATE: 1997-10-24

PRIOR PLING DATE: 1997-10-24

PRIOR PELING DATE: 1997-10-24

PRIOR PILING DATE: 1997-10-28

PRIOR APPLICATION NUMBER: 60/063540

PRIOR APPLICATION NUMBER: 60/063541

PRIOR PILING DATE: 1997-10-28

PRIOR PILING DATE: 1997-10-28

PRIOR PILING DATE: 1997-10-28

PRIOR APPLICATION NUMBER: 60/063544

PRIOR APPLICATION NUMBER: 60/063544
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Matches:
Conservative:
Mismatches:
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2245.50
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ORGANISM: Homo Sapien
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APPLICANT: Chen, Jian
APPLICANT: Chen, Jian
APPLICANT: Chen, Jian
APPLICANT: Coddard, Audrey
APPLICANT: Godowski, Paul J.
APPLICANT: Gorney, Austin L.
APPLICANT: Ban, James
APPLICANT: Pan, James
APPLICANT: Pang, Zemin
TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: P3430R16519
CURRENT PPLICATION NUMBER: 10/052586
PRIOR APPLICATION NUMBER: 10/052586
PRIOR APPLICATION NUMBER: 60/059263
PRIOR FILING DATE: 2002-01-15
PRIOR FILING DATE: 1997-09-18
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AATGATCCAAAGTTCATTAGTGCCCACCTCATCTCAGAGGGGGGAGTACCTGAAGATGAC 922
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Sequence 348, Application US/10208024
Publication No US20040048335A1
GENERAL INFORMATION:
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Watanabe, Colin K.
Wood, William I.
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Gurney, Austin L.
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LeuGlyProThrHisAspHisHisTyrIleArgThrAspIleSerGluHisTyrTrpLeu
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PRIOR APPLICATION NUMBER: 60/063486

PRIOR FILING DATE: 1997-10-21

PRIOR PELOR FILING DATE: 1997-10-21

PRIOR APPLICATION NUMBER: 60/063541

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PRIOR APPLICATION NUMBER: 60/063544

PRIOR APPLICATION NUMBER: 60/063544

PRIOR PILING DATE: 1997-10-28

PRIOR FILING DATE: 1997-10-28

PRIOR PILING DATE: 1997-10-28

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Mismatches:
Indels:
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2060 ATCAGAGTGGATGATCATATCATCAGGACAGATCAAGGCCTTCTGCTACGTAGTCTACAA 2119
                                                                                         CAGAAGGATTCAGGCAATTAGCTCTGCCATGCGTGGAACATGGGTTCATACAAACTCTT 2179
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                      748 ------ProLysTrpLysHisMetGlnGluMetLysLysLysArgAsnArgArg 763
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TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: P943-0017-04-65
CURRENT APPLICATION NUMBER: 105/10/201,853
CURRENT FILING DATE: 2002-07-23
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PRIOR PUBLICATION NUMBER: 10/052566

PRIOR PAPLICATION NUMBER: 0/052566

PRIOR FILING DATE: 12097-09-18

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PRIOR APPLICATION NUMBER: 60/063121

PRIOR PILING DATE: 1997-10-24

PRIOR APPLICATION NUMBER: 60/063486

PRIOR PILING DATE: 1997-10-21

PRIOR PILING DATE: 1997-10-28

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GENERAL INFORMATION:
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Smith, Victoria
Watanabe, Colin K.
Wood, William I.
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Godowski, Paul J.
Gurney, Austin L.
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179 GlyArgLeuLysCysProPheAspProGlnGlnProPheAlaSerValMetThrAspGlu 198
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; Prior Application data removed - See File Wrapper or PALM.; NUMBER OF SEQ ID NOS: 612; SEQ ID NO 348; LENGTH: 777
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Oy 2180 CTTAAGGTAACCCTGGAAGTCATTGACACAGAGCAI	2360 730 2420 748	<pre>Qy 2480 ACCCACGAATTTGAGAGGCACCCAGGAGTGTC</pre>	APPLICANT: Chen, Jian APPLICANT: Desnoyers, Luc APPLICANT: Goddard, Audrey APPLICANT: Goddard, Audrey APPLICANT: Gurney, Austin L. APPLICANT: Pan, James APPLICANT: Rath, Victoria APPLICANT: Matenabe, Colin K. APPLICANT: Watenabe, Colin K. APPLICANT: Applicant: Zhanq, Zemin	TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE TITLE OF INVENTION: ACIDS ENCODING THE SAME FILE REPERBENCE: P3430R1C41 CURRENT APPLICATION NUMBER: US/10/174,581 CURRENT FILING DATE: 2002-06-18 FRIOR APPLICATION NUMBER: 06/052586 PRIOR FILING DATE: 2002-01-15 PRIOR FILING DATE: 1997-09-18 PRIOR APPLICATION NUMBER: 60/05266 PRIOR APPLICATION NUMBER: 60/05266 PRIOR APPLICATION NUMBER: 06/05250	PRIOR FILING DATE: 1997-10-17 PRIOR APPLICATION NUMBER: 60/063120 PRIOR PELLING DATE: 1997-10-24 PRIOR APPLICATION NUMBER: 60/063121 PRIOR PELLING DATE: 1997-10-24 PRIOR FILING DATE: 1997-10-24 PRIOR FILING DATE: 1997-10-24 PRIOR PELLING DATE: 1997-10-28 PRIOR PELLING DATE: 1997-10-28 PRIOR FILING DATE: 1997-10-28 PRIOR FILING DATE: 1997-10-28	PRIOR APPLICATION NUMBER: 60/063544 PRIOR FILING DATE: 1997-10-28 PRIOR PILING DATE: 1997-10-28 PRIOR PELLING DATE: 1997-10-28 PRIOR PELLING DATE: 1997-10-29 PRIOR PELLING DATE: 1997-10-29 PRIOR PLLING DATE: 1997-10-29 PRIOR PLLING DATE: 1997-10-31 PRIOR APPLICATION NUMBER: 60/063870 PRIOR PELLING DATE: 1997-10-31 PRIOR FILING DATE: 1997-10-31
GACACTCATTTTGATGAACTGCAGGATGTATTCCTAATGAACTTTAAAGATCCTAAAAAT	359 ValTyrSerMetAlaAspileArgAlaValPheAsnGlybroTyrAlaHisbysGluSer 378 1283 CCCAACTATCAATGGGTGCCTTATCAAGGAAGAGTCCCCTATCACGGCCAGGAACTTGT 1342 1379 AlaAspHisArgTrpValGlnTyrAspGlyArgIleProTyrProArgProGlyThrCys 398 1343 CCCAGGAAACATTTGGTGGTTTTGACTACAAAGGACCTTGATGATTATA 1399 11	1400 ACCTITGCAAGAAGTCATCCAGCCATGTACAATCCAGTGTTTCCTATGACCAATG5 ::: :::		1700 CIATATATTGGTTCAACGGCTGGGGTTGCCCAGCTCCCTTTACACCGGTGTGATATTTAC 1759	1880 GGAGACCCACTGACTCACACTTACACCATGATAATCACCATGGCCACAGCCCT 1939 [:::	617 SerGlnGlnalaThrIleLysTrpTyrIleĠinÀrgSerGlyAspĠiuHisArgĠiuĠiu 636 2060 ATCACAGTGGATGATCATCATCAGGACAGATCAAGGCTTTCTGCTACGTACG
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tgGlnArgAsnLysGlyGly----- 747 AAAATAAGAAAGGTAGAAACAGGAGG 2479 || ||||||| |UwetLysLysLysArgAsnArgAgag 763 AGCATTTGGAAGAACTTCTTCATAAAGAT 2239 CCAATAGCATGACACCTAGCCAGAAG 2299 ACCCCAATCTCAACACGATGGATGAG 2359 :::||| |InMetGluAsnThrGlnArgAlaGlu 696 ::: ----AspLeuLeuAlaGluSerArg 710 E POLYPEPTIDES AND NUCLEIC TC 2512 || || 775

PRIOR APPLICATION NUMBER: 60/066131
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PRIOR FILING DATE: 1998-05-15
PRIOR PRIOR OFFICATION NUMBER: 60/08573
PRIOR PAPLICATION NUMBER: 60/08579
PRIOR PALLING DATE: 1998-05-15
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PRIOR PALICATION NUMBER: 60/08025
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PRIOR APPLICATION NUMBER: 60/089514
PRIOR FILING DATE: 1998-06-16
PRIOR APPLICATION NUMBER: 60/089538
PRIOR FILING DATE: 1998-06-17
PRIOR APPLICATION NUMBER: 60/089598
PRIOR FILING DATE: 1998-06-17
PRIOR APPLICATION NUMBER: 60/089653
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APPLICANT: Goddard, Audrey
APPLICANT: Goddard, Audrey
APPLICANT: Goddard, Paul J.
APPLICANT: Goddard, Paul J.
APPLICANT: Pan, James
APPLICANT: Pan, Vancoria
APPLICANT: Watanabe, Colin K.
APPLICANT: Wood, William I.
APPLICANT: Wood, William I.
TILE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
TILLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REPERBRY APPLICATION NUMBER: US/10/176,483
CURRENT APPLICATION NUMBER: US/202-06-20
Prior application removed - See File Wrapper or Palm
NUMBER OF SEQ ID NOS: 612
2180 CITAAGGIAACCCTGGAAGTCATTGACACAGAGCATTGGAAGAACTTCTTCATAAAGAT 2239
                                                     2240 GATGATGGAGATGGCTCTAAGACCAAAGAATGTCCAATAGCATGACACCTAGCCAGAAG 2299
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                                                                                                                                                                                                                                                                                                                                          Sequence 348, Application US/10176483
Publication No. US20030017541A1
GENERAL INFORMATION:
APPLICANT: Baker Kevin P.
APPLICANT: Chen, Jian
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Godowski, Paul J.
Gurney, Austin L.
Pan, James
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Baker, Kevin P.
Chen, Jian
Desnoyers, Luc
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; ORGANISM: Homo Sapien
US-10-176-749-348
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Best Local Similarity:
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| GlnilePheLysHisSerSerllelleLeuAsnMetGluLeuSerLeuLysGlnGlnGln
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| GlyAspProlleThrGlnCysTrpAspIleGluAspSerIleSerHis---GluThrAla
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 GCAGAAGATGGACAGTATGATGTTATGTTTATCGGAACAGATGTTGGGACCGTTCTTAAA
             AlaCysSerArgTyrAlaProThrSerLysArgArgAlaArgArgGlnAspValLysTyr
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Sequence 348, Application US/10176749 Publication No. US20030017542A1

APPLICANT: Goddard, Audrey
APPLICANT: Goddard, Audrey
APPLICANT: Goddard, Audrey
APPLICANT: Goddwski, Paul J.
APPLICANT: Gurney, Austin L.
APPLICANT: Pan, James
APPLICANT: Pan, James
APPLICANT: Pan, James
APPLICANT: Wacanabe, Colin K.
APPLICANT: Wacanabe, Colin K.
APPLICANT: Zhang, Zemin
TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
TITLE OF INVENTION: ACIDS ENCODING THE SAME,
FILE REFERENCE: P34.90R.C76
CURRENT PILING DATE: 2002-06-20
CURRENT FILING DATE: 2002-06-20
Prior application removed - See File Wrapper or Palm
NUMBER OF SEQ ID NOS: 612
SERVINE: 777 455 ---AAGGATTTTCAAAAGATTGTGTGGCCAGTATCTTACACCAGAAGAGATGAATGCAAG 511 119 LeuAlaGlyLysAspAlaAsnThrGluCysAlaAsnPheIleArgValLeuGlnProTyr 138 199 TyrLeuTyrSerGlyThrAlaSerAspPheLeuGlyLysAspThrAlaPheThrArgSer 218 394 631 GAAATTGGACATCATCCTGAGGACAATATTTTTAAGCTGGAGAACTCACATTTTGAAAAC 691 862 AAGAACAATGTGCCAAGGCTGAAATTATCCTACAAAGAAATGTTGGAATCCAACAATGTG 334 CGGAGTAGGCTGTATGTTGGAGCAAAGGATCACATATTTTCATTCGACCTGGTTAATATC 454 GGCCGTGGGAAGAGTCCATATGACCCTAAGCTGCTGACAGCATCCCTTTTAATAGATGGA 751 512 IGGGCIGGAAAAGACAICCTGAAAGAAIGIGCIAAITICAICAAGGIACTIAAGGCAIAI AATCAGACTCACTTGTACGCCTGTGGAACGGGGGCTTTTCATCCAATTTGCACCTACATT GAATTATACTCTGGAACTGCAGCTGATTTTATGGGGCGAGACTTTGCTATCTTCCGAACT 777 413 141 177 21 Matches: Conservative: Mismatches: Indels: US-09-774-490-1 (1-2709) x US-10-176-749-348 (1-777) ength: 1.9e-201 2245.50 73.678 54.928 45.868 812

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                                                           863 AATGATCCAAAGTTCATTAGTGCCCACCTCATCTCAGAGAGTGACAATCCTGAAGATGAC 922
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               697 HisGluGluGlyGlnValLys-------AspLeuLeuAlaGluSerArg 710
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
US-09-653-274-4
US-09-653-274-13
US-09-653-274-13
US-09-653-274-13
US-09-06-052-62
US-09-06-060-62
US-09-06-060-62
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US-09-254-594-6
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US-08-121-113D-58
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NAME: Osman, Richard A.
REGISTRATION NUMBER: 36.627
REFERENCE/DOCKET NUMBER: B94-002-1
TELECOMMUNICATION INFORMATION:
TELEPRA: (415) 343-4342
TELEFAX: (415) 343-4342
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FILING DATE: 13-SEP-1993
CLASSIFICATION: 514
    ZIP: 94104
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MODEL=frame+ n2p.model -DEV=xlp

-Q=/cgn2 1/USBTO spool p/USO974490/runat 21092004 144246 2377/app query.fasta_1.2887

-Q=/cgn2 1/USBTO spool p/USO974490/runat 21092004 144246 2377/app query.fasta_1.2887

-DB=ISSUEd Patents AA -OFMT=fastan -SUFFIX=rai -MINMATCH=0.1 -LOOPCL=0

-LOOPEXT=0 -UNITS=Bits -START=1 -END=-1 -MARTIX=blosum62 -TRANS=human40.cdi
-LIST=45 -DOCALIGN=200 -THR SCORE=pct -THR MAX=100 -THR MIN=0 -ALIGN=15

-MODE=LOCAL -OUTFWT=pco -NORM=ext -HARDSIZE=500 -MINLEN=0 -MAXIEN=200000000

-USER=USO9774490 @CGN 1 1.74 @runat 21092004 144246 2377 -NCPU=6 -ICPU=3

-NO MMAP -LARGEQUERY -NEG SCORES=0 -WAIT -DSPBLOCK=100 -LONGLOG

-DEV TINBOUT=120 -WARN TIMEOUT=30 -THRRADS=1 -XGAPOP=10 -XGAPEXT=0.5 -FGAPOP=6

-FGAFEXT=7 -YGAPOP=10 -YGAPEXT=0.5 -DELOP=6 -DELEXT=7
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1: /GGT2_6/ptodata/2/iaa/5A_COMB.pep:*
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3: /CGT2_6/ptodata/2/iaa/6A_COMB.pep:*
4: /CGT2_6/ptodata/2/iaa/6B_COMB.pep:*
5: /CGT2_6/ptodata/2/iaa/FGTCGCOMB.pep:*
6: /CGT2_6/ptodata/2/iaa/FGTCGCOMB.pep:*
                      GenCore version 5.1.6
(c) 1993 - 2004 Compugen Ltd.
                                                                                                       - protein search, using frame_plus_n2p model
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INFORMATION FOR SEQ ID NO: 19 SEQUENCE CHARACTERISTICS: LENGTH: 771 amino acids TYPE: amino acids
                              ; TYPE: amino acid
; TOPOLOGY: linear
; MOLEGULE TYPE: protein
US-08-121-713D-54
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1040 AATAAATGGACAACATTCCTCAAAGCTCGTCTGATTTGCTCAGTGCCAGGTCCAAATGGC 1099
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        181 LeulleAspGlyGluLeuTyrSerGlyThrAlaAlaAspPheMetGlyArgAspPheAla 200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GAATCCAACAATGTGATCACTTTCAATGGCTTGGCCAACAGCTCCAGTTATCATACCTTC 379
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Matches:
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Query Match:
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GATGATGGAGATGGCTCTAAGACCAAAGAAATGTCCAATAGCATGACACCTAGCCAGAAG
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                                                                                              CTTAAGGTAACCCTGGAAGTCATTGACACAGAGCATTTGGAAGAACTTCTTCATAAAGAT
                                                                                                                         LeuLysValThrLeuGluValIleAspThrGluHisLeuGluGluLeuLeuHisLysAsp
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GENERAL INFORMATION:
APPLICANT: KOLOGKIN, Alex L.
APPLICANT: KOLOGKIN, Alex L.
APPLICANT: Matthes, David
APPLICANT: Matthes, David
APPLICANT: Matthes, David
APPLICANT: DATENCY TIMOTHY
TITLE DF INVENTION: The Semaphorin Gene Family
NUMBER OP SEQUENCES: 100
CORRESPONDENCE ADDRESS:
ADDRESSEE: SCIENCE & TECHNOLOGY LAW GROUP
STREET: 2GB Bush Street, Suite 3200
CITY: San Francisco
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Patent No. 5807826
GENERAL INFORMATION:
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SEQUENCE CHARACTERISTICS:
LENGTH: 771 amino acids
TYPE: amino acid
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TOPOLOGY: linear
MOLECULE TYPE: protein
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Db 661 LeuLysValThrLeuGluValIleAspThrGluHisLeuGluGluLeuLeuHisLysAsp 680 Qy 2240 GATGATGGAGATGGCTCTAAGACCAAAAGAAATGTCCAATAGCATGACACCAGCAGAAG 2299		2300 GTCTGGTACAGAGTTCATGCAGCTCATCAACCACCCCAATCTCAACAGGATGGAT	Oy 2360 TICTGTGAACAGTTTGGAAAAGGGACCGAAAACAACGTCGGCAAAGGGCAGGACATACC 2419	Oy 2420 CCAGGGAACAGTAACAAATGGAAGCACTTACAAGAAATAAGAAAGGTAGAAACAGGAGG 2479	Oy 2480 ACCCAGGAATTTGAGGGCACCCAGGAGTGTC 2512	NESULT 3 VS-03-54 ; Sequence 54, Application US/09060692 ; Patent No. 5935865	; GENERAL INFORMATION: ; APPLICANT: GOOdman, COTEY S. ; APPLICANT: Kolodkin, Alex L. ; APPLICANT: Matthes, David	APPLICANT: Bentley, David R. APPLICANT: O'Connor Timothy ITITE OF INVENTION: The Semaphorin Gene Family NUMBER OF SEQUENCES: 100	CORRESPONDENCE ADDRESS: ADDRESSEE: SCIENCE & TECHNOLOGY LAW GROUP STRRET: 268 Bush Street, Suite 3200 CITY: San Francisco	COUNTRY: USA ZIP: 94104 COMPUTER READABLE FORM:) MEDIUM TYPE: Floppy disk ; COMPUTER: PLOPDY disk ; OPBRATING SYSTEM: PC-DOS/MS-DOS ; SOFTWARE: Patentin Release #1.0, Version #1.25	CURRENT APPLICATION DATA: APPLICATION NUMBER: US/09/060,692 FILING DATE: CLASSIFICATION: 514	; PRIOR APPLICATION DATA: ; APPLICATION NUMBER: US/08/121,713 ; FILING DATE: 13-SEP-1993 ; ATTORNEY/AGENT INFORMATION:	A S S A	; TELEPHONE: (415)343-4341 ; TELEAX: (415)343-4342 ; TELEX: : INFORMATION FOR SEO ID NO: 54:	CHARACTERISTICS: 771 amino acids amino acid II. linear) MOLECULE TYPE: protein US-09-060-692-54	Pred. No.: 0 Length: 771 Score: 4201.00 Matches: 771 Percent Similarity: 100.00% Conservative: 0
	1160 AATCCAGTTGTATATGGAGTGTTTACGACTTCCAGTAACATTTTCAAGGGATCAGCCGTG 1219 	1220 TGTATGTATAGCATGAGTGATGTGAGAAGGGTGTTCCTTGGTCCATATGCCCACAGGGT 1279	1280 GGACCCAACTATCAATGGGTGCCTTATCAAGGAAGAGTCCCCTATCCAGGCAAGGAACT 1339 	ATGTTATA	ACCTITGGAAGAAGTCATCGAGCCATGTACAATCCAGTGTTTCCTATGAACAACAACGCCCA 	1460 ATAĞTGATCAAAACGGANGTAAATTATCAATTTACACAAATTGTCGTAGACCGAGTGGAT 1519 	1520 GCAGAAGATGGACAGTATGATGATTTATCGGAACAGATGTTGGGACCGTTCTTAAA 1579 	1580 GTAGTTTCAATTCCTAAGGAGACTTGGTATGATTTAGAAGAGGTTCTGCTGGAAGAATG 1639 	1640. ACAGITTITCGGGAACCGACTGCTATTTCAGCAATGGAGCTTTCCACTAAGCAGCAACAA 1699 	1700 CTATATATTGGTTCAACGGCTGGGGTTGCCCAGCTCCCTTTACACCGGTGTGATATTTAC 1759	1760 GGGAAAGGGTGTGCTGAGTGTTGCCTGGCCCGGAGACCCTTACTGTGCTTGGGATGGTTCT 1819	1820 GCATGTTCTCGCTATTTTCCCACTGCAAAGAGACGCACAAGACGACAAGATATAAGAAAT 1879 	1880 GGAGACCCACTGACTGTTCAGACTTACACCATGATAATCACCATGGCCCACAGCCCT 1939	1940 GAAGAGAGAATCATCTATGGTGTAGAGAATAGTAGCACATTTTTGGAATGCAGTCCGAAG 1999 	2000 TCGCAGAGAGCGCTGGTCTATTGGCAATTCCAGAGGGCGAAATGAAGAGCGAAAAGAAGAG 2059 	ATCAGAGTGGATGATCATATCAGGACAGATCAAGGCCTTCTGCTACCAAGGTTACAAA	2120 CAGAAGGATTCAGGCAATTACCTCTGCCATGGGTGGAACATGGGTTCATACAACTCTT 2179	CTTAAGGTAACCCTGGAAGTCATTGACACAGAGCATTTGGAAGAACTTCTTCATAAAGAT
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GTCTGGTACAGAGACTTCATGCAGCTCATCAACCACCCCAATCTCAACACGATGGATG	ACCACGARTTGAGAGGGCACCAGAGTGTC 2512	SULT 4 -08-833-391-54 Sequence 54, Application US/08833391 Patent No. 6013781 GENERAL INFORMATION:	APPLICANT: GOOdman, Cotey's. APPLICANT: Matches, David APPLICANT: Bentley, David R. APPLICANT: Bentley, David R. APPLICANT: Goodwan, Thoughout, Good Family.	SECURIOS: 100 BENCE ADDRESS: ES SCIENCE & TECHNOLOGY LAW GROUP	SYMEN: Zes Bush Street, Suite 3200 STATE: CA STATE: CA TEN. 04.04	COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk COMPUTER: IBM PC compatible COMPUTER: TO PC-DOS/MS-DOS	SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/833,391	CLASSITION: 530 PRIOR APPLICATION: 530 PRIOR APPLICATION DATA: PAPLICATION NUMBER: US 08/121,713	ATTORNEY/AGENT INCOMMATION: NAME: Osman, Richard A. REGISTRATION NUMBER: 36.27 PERPERENCE / DOCUMER: B94-002-1	LECOMMUNICATION INFORMATION: TELEPHONE: (415)343-4341 meney: (415) 343-4342	INFORMATION FOR SEQ ID NO: 54: SEQUENCE CHARACTERISTICS: LENGTH: 771 amino acids	11F5: definition actu MOLECULE TYPE: protein 8-833-391-54	0 4201.00 100.00%	Best Local Similarity: 100.00% Mismatches: 0 Query Match: 85.80% Indels: 0 DB: 3 Gaps: 0	US-09-774-490-1 (1-2709) x US-08-833-391-54 (1-771)

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721 PheCysGluGlnValTrpLysArgAspArgLysGlnArgArgGlnArgProGlyHisThr 740
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                                     CCAGGGAACAGTAACAAATGGAAGCACTTACAAGAAAATAAGAAAGGTAGAAACAGGAGG
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; Sequence 54, Application US/0966610
; Sequence 54, Application US/0966610
; Patent No. 6344544
; GENERAL INFORMATION:
; APPLICANT: Goodman, Corey S.
APPLICANT: Matthes, David R.
APPLICANT: Matthes, David R.
APPLICANT: Bentley, David R.
APPLICANT: David R.
APPLICANT: David R.
APPLICANT: Hooky, David R.
APPLICANT: David R.
APPLICANT: Sequences: 100
CORRESPONDENCES: 100
CORRESPONDENCE ADDRESS:
ADDRESSER: SCIENCE FIECHNOLOGY LAW GROUP
                                                                                                                                                                                                                                                                                                                                                                                                                   CITY: San Francisco
CITY: San Francisco
COUNTRY: USA
ZIP: 94104
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM For Compactible
COMPUTER: TRANCE COMPACTIBLE
COMPATION SYSTEM: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/060,610
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Conservative:
Mismatches:
Indels:
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CLASSIFICATION:
PRIOR PEPLICATION DATA:
PRIOR PEPLICATION DATA:
APPLICATION DATE:
ATTORNEY/AGBNT INFORMATION:
NAME: Osman, Richard A.
REGISTRATION NUMBER: 36 627
REFERENCE/DOCKET NUMBER: B94-002-TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 343-4341
TELEPAX: (415) 343-4342
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SEQUENCE CHARACTERISTICS:
LENGTH: 771 amino acid
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Best Local Similarity:
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       GATGAATGCAAGTGGGCTGGAAAAGACATCCTGAAAGAATGTGCTAATTTCATCAAGGTA
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                                                                                                                                         TGCACCTACATTGAAATTGGACATCATCCTGAGGACAATATTTTTAAGCTGGAGAACTCA
                                                                                                                                                             CysThrTyr1leGlu1leGlyHisHisProGluAspAsnIlePheLysLeuGluAsnSer
                                                                                                                                                                                                       CATITIGAAAACGGCCGTGGGAAGAGTCCATATGACCCTAAGCTGCTGACAGCATCCCTT
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                            AspGluCysLysTrpAlaGlyLysAspIleLeuLysGluCysAlaAsnPheIleLysVal
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                                                                     CTTAAGGCATATAATCAGACTCACTTGTACGCCTGTGGAACGGGGGCTTTTCATCCAATT
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                                                                                                Sequence 54, Application PC/TUS9410151A
GENERAL INFORMATION:
APPLICANT: The Regents of the University of California
TITLE OF INVENTION: The Semaphorin Gene Family
NUMBER OF SEQUENCES: 66
CORRESPONDENCE ADDRESS:
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MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC compatible
COMPUTER: IEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US94/10151A
FILING DATE: 13-SEP-1994
                                                                                                                                                                                                ADDRESSEE: FLEHR HOHBACH TEST ALBRITTON & HERBERT STREET: 4 Embarcadero Center, Suite 3400 STREET: San Francisco STATE: CA COUNTRY: USA
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Matches:
Conservative:
Mismatches:
Indels:
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NAME: Osman, Richard A.
REGISTRATION NUMBER: 36,627
REFERENCE/DOCKET NUMBER: FP-58750-PC/RAO
TELEPHONICATTON INFORMATION:
TELEPHONE: (415) 789-3249
TELEFRAX: (415) 398-3249
TELEFRAX: (415) 398-3249
INFORMATION FOR SEQ ID NO: 554:
SEQUENCE CHARACTERISTICS:
LENGRATION FOR SEQ ID NO: 54:
TYPE: amino acids
TYPE: amino acids
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TYPE: 1 mino acids
TYPE: 1 mino acids
TYPE: 1 mino acids
TYPE: 1 mino acids
TOPOLOGY: 1 linear
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Best Local Similarity:
Query Match:
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Sequence 3, Application US/08556422A
Patent No. 6576754
GENERAL INFORMATION:
APPLICANT: HALL, Kathryn T.
APPLICANT: FREEMAN, Gordon J.
APPLICANT: SCHULTEZ, Joachim L.
APPLICANT: BOUSSIOTIS, Wassiliki
APPLICANT: NADLER, Lee M.
US-08-556-422A-3
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241 AsplysValTyrPhePhePheArgGluAsnAlaIleAspGlyGluHisSerGlyLysAla 260
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TITLE OF INVENTION: NUCLEIC ACIDS ENCODING CD100 MOLECULES FILE REFERENCE: DFN-005CPA2 CURRENT APPLICATION NUMBER: US/08/556,422A CURRENT FILING DATE: 1995-11-09 NUMBER OF SEQ ID NOS: 7 SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 3 LENGTH: 655
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Qy 1019 GGAGGGCACAGAAGTCTGGTGAATAAATGGACJ 	2000 TCGCAGAGAGCGCTGGTCTATTGGCAATTCCAGAGGCGAAATGAAGAGCGAAAAGAGAGAG
Qy 959 GGAGAACACTCTGGAAAGCTACTCACGCTAGI	1940 GAAGAGAATCATCTATCGTGTAGAAATAGTAGCACATTTTTGGAATGCAGTCCGAAG 1999
Qy 899 GAGAGTGACATCCTGAAGATGACAAGTATA(GGAGACCCACTGACTCAC
-09-774-490-1 (1-2709) x US-08-136-92	1820 GCATGTTCTCGCTATTTTCCCACTGCAAAGACGACGACAAGACGACAAGATATAAGAAAT 1879
L Similarity: 99.37% ocal Similarity: 99.37% March: 1	GGGAAAGCGTGTGCTGAG'
Alignment Scores: 2.31e-246 Length:	1700 CIAIATATTAGITCAACGGCIGGGGTIGCCCGGCTCCCTTTACACGGGTGTGATTTAC 1759
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; SEQUENCE CHARACTERISTICS; ; LENGTH: 477 amino acids . TVDF: smino acids	.*o. valvaiseillekrolybsillinrirpiyraspleudiudiuvaileuleudiuMet 480 1640 ACAGTTTTCGGGAACCGACTGCTATTCAGCAATGGAGGTTTCCACTAAGCAAGAA 1699
TELEFAX: 215-568-3419 ; INFORMATION FOR SEQ ID NO: 2:	GTAGTTTCAATTCCTAAGGAGACTTGGTATGATTAGAAGAGGTTCTGCTGGAAGAAATG
REGISTRATION NUMBER: 33,229 REFERENCE/DOCKET NUMBER: UPN-1428 TELECOMMUNICATION INFORMATION:	1520 GCAGAAGATGGATGATGATGATTATGGGAACAGATGTTGGGACGGTTGTAAA 1579
CLASSIFICATION: 435 ATTORNEY AGENT INFORMATION: NAWE: Deluca, Mark	421 IleValIleLysThrAsp
; APPLICATION NUMBER: US/08/136,922 ; FILING DATE:	1460 ATAGTGATCAAAAGGGATGTAAATTATCAAAATTTGTCGTAGACGGAGTGGAT 1519
25.25	1400 ACCTTTGCAAGAAGTCATCCAGCCATGTACAATCCAGGTTTTCCTATGAACAATCGCCCA 1459
COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk	1340 TGTCCCAGCAAACATTTGGTGGTTTTGACTCTACAAAGGACCTTCCTGATGATGTTATA 1399
; STREET: One Liberty Place ; CITY: Philadelphia ; STATE: PA . COUNTRY PLACE	1280 GGACCCAATTATCAATGGGTGCCTTATCAAGGAAGAGTCCCCTATCCACGGCCAGGAACT 1339
NUMBER OF SECUENCES: 2 CORRESPONDENCE ADDRESS: ADDRESSE: ADDRESSE: Moodcock Mashburn Kurtz M	341 CysMetTyrSerMetSerAspValArgArgValPheLeuGlyProTyrAlaHisArgAsp 360
APPLICANT: Raper, Jonathan A. APPLICANT: Luo, Yuling ITLLE OF INVENTION: Compositions Which R. TITIS OF INVENTION: Receneshation and Mark	AATCCAGTTGTATATGGA
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                                       ATGA CTTTAAAGATCCTAAAATCCAGTTGTATATGGAGTGTTTACGACTTCCAGTAAC
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422 HisGlyPhelleGlnThrLeuLeuLysValThrLeuGluValIleAspAsnGluHisLeu 441
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Mismatches:
Indels:
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APPLICANT: INAGAKI, Shinobu
APPLICANT: INAGAKI, Shinobu
ITILE OF INVENTION: NOVEL SEMAPHORIN GENES (I)
ITILE OF INVENTION: NOVEL SEMAPHORIN GENES (I)
FILE REFERENCE: 0020-4562P
CURRENT PELLING DATE: 1999-05-14
PRIOR PELLING DATE: 1997-11-12
PRIOR FILING DATE: 1997-11-12
PRIOR FILING DATE: 1996-11-15
NUMBER OF SEQ ID NOS: 4
SOFTWARE: PATENTING DATE: 1996-11-15
NUMBER OF SEQ ID NOS: 4
SOFTWARE: PATENTING DATE: 1996-11-15
NUMBER OF SEQ ID NOS: 4
SEQ ID NO 1
ERNGTH: 775
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Matches:
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Best Local Similarity:
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127 ArgThrHisLeuLeuThrCysAlaThrGlyAlaPheAspProHisCysAlaPheIleArg 146

575 CAGACTCACTTGTACGCCTGTGGAACGGGGGTTTTCATCCAATTTGCACCTACATTGAA 634

635 ATTGGACATCTTCTGAGGACAATATTTTAAGCTGGAGAACTCACATTTTGAAACGGC 694

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TYPE: PRT
ORGANISM: Homo sapiens
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US-08-556-422A-2
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ValGlyHisHisSerGluGluProLeuPheHisLeuGluSerHisArgSerGluArgGly 166
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LeuPhcAlaGlyLeuTyrSerAspTyrTrpGlyArgAspSerAlaIlePheArgSerMet
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                                                  TTATACTCTGGAACTGCAGCTGATTTTATGGGGCGAGACTTTGCTATCTTCCGAACTCTT
                                                                                    GGGCACCACCACCAATCAGACAGAGCAGCATGATTCCAGGTGGCTCAATGATCCAAAG
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|ThrAspGlyLysTyrAsnLeuArgGlnLeuAlaValAspArgValGluAlaGluAspGly
                 CGTGGGAAGAGTCCATATGACCCTAAGCTGCTGACAGCATCCCTTTTAATAGATGGAGAA
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AsnalaalaGlnGlnCysPheGlyGlnGlnPheValGlyAspAlaLeuAspArgThrĠiu
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LysThrAspAspArgValValLysMetAspLeuGlyLeuLeuPheLeuArgValArgLys
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                             AlaAspCysCysLeuAlaArgAspProTyrCysAlaTrpAspGlyIleSerCysSerArg
                                                                                                TATTTTCCCACT-----GCAAAGAGACGCACAAGAGACGACAAGATATAAGAAATGGA
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Patent No. 657674
GENERAL INFORMATION:
APPLICANT: HALL, Kathryn T.
APPLICANT: RREMAN, Gordon J.
APPLICANT: SCHULTZE, Joachim L.
APPLICANT: BRUSSIOTIS, Vassiliki
APPLICANT: BRUSSIOTIS, Vassiliki
APPLICANT: NADLER, Lee M.
TITLE OF INVENTION: NUCLEIC ACIDS ENCODING CD100 MOLECULES
FILE REFREENCE: DRA-0856PA2
CURRENT APPLICATION NUMBER: US/08/556,422A
CURRENT RILING DATE: 1995-11-09
NUMBER OF SEQ ID NOS: 7
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 2
IENGTH: 862
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us-09-774-490-1.rai

US-08-55	6-422A-2
Alignmen Pred. No Score: Percent Best Loc Query Mai	Scores: 7.83e-89 Length: 862 Similarity: 51.42% Conservative: 115 al Similarity: 35.04% Mismatches: 218 tch: 4 Gaps: 26
US-09-77	4-490-1 (1-2709) x US-08-556-422A-2 (1-862)
දු දු	284 GTGCCAAGGCTGAAATTATCCTACAAAGAAATGTTGGAATCCAACAATGTGATGACACTTTC 343 ::: ::: 27 IleProArglieThrTrpGluHisArgGluValHisLeuValGluPhe 42
ζŏ	4 AATGGCTTGGCCAACAGCTCCAGTTATCATACCTTCCTTTTGGATGAGGAACGGAGTAGG 403
qq	43 HisGluProAspIleTyrAsnTyrSerAlaLeuLeuLeuSerGluAspLysAspThr 61
දු ද	404 CIGIAIGITGGAGCAAAGGAICACATATTTCATTCGACCIGGTTAATATCAAGGATTTT 463 Db
3 &	2 Deutyliteolynianlystuniavalruchavalnishinanaruhniiteocituuys oi 4 CAAAAGATTGTGTGGCCAGTATCTTACACCAGAAGAGAAG
QQ	82 GlnHisGluValTyrTrpLysValSerGluAspLysLysAlaLysCysAlaGluLysGly 101
රු ද	ACATCCTGAAAGAATGTGCTAATTTCATCAAGGTACTTAAGGCATATAATCAGACT 580 **Interpretation of the content of
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Dp	ThrSerPheLysPheLeuGlyLysAsnGluAspGlyLysGly 154
A S	701 AAGAGTCCATATGACCTAAGCTGACGCTGACCCCTTTTAATAGATGGAGAATTATAC 760 ::: 155 ArgCysProPheAspProAlaHisSerTyrThrSerValMetValAspGlyGluLeuTyr 174
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<u>ن</u> د	821 CACCACCCAATCAGGACAGAGCAGCATCCAGGGGCTCAATGATCCAAAGTTCATT 880 104 History Computer of the co
ì à	81 AGTGCCCACCTCATCTCAGAGAGTGACAATCCTGAAGATGACAAAGTATAC 931
- q _Q	::: ProAspSerProAspGlyGluAspAspArgValTyr 232
٥x	932 ITITICITCGGGAAAAGGAAGAAGGAAAAGCTACGCGAGA 991 DD
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දු දු	992 ATAGGTCAGATATGCAGAATGACTTTGGAGGCACAGAAGTCTGGTGAATAAATGGACA 1051
Š	1052 ACATTCCTCAAAGCTCGATTTGCTCAGGGCCCAAAATGGCATTGACACTCAT 1111
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& 8	1112 TITGATGAACTGCAGGATGTATTCCTAATGAACTITAAAGATCCTAAAAAATCCAGTTGTA 1171
ò	72 TATGGAGTGTTTACGACTTCCAGTAACATTTTCAAGGGATCAGCCGTGTGTATGTA

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123;	2 ATGAGTGATGTGAGAGAGGTGTTCTTGGTCCATATGCCCACAGGGAT 1279 :::
350	0 GGACCCAACTATCAATGGGTGCCTTATCAAGGAAGGTCCCCTATCCACGGCCAGGAACT 1339 :::::::::::::::::::::::::::::::::::
134	0 TGTCCCAGCAAAACATTTGGTGGTTTTTGACTCTACAAAGGACCTTCCTGATGAT 1393 ::: 0 CysileAspSerGluAlaAsgAlaAsnTyrThrSerSerbeuAsnLeuPrOAspLys 389
139,	4 GTTATAACCTTTGCAAGAAGTCATCCAGCCATGTACAATCCAGTGTTTCCTATGAACAAT 1453 :::
145,	4 CGCCCAATAGTGATCAAAACGGATGTAAAATTATCAATTTACACAAATTGTCGTAGACCGA 1513
1514	4 GTGGATGCAGAAGATGGACAGTATGATGTTATGTTTATCGGAACAGATGTTGGGACC 1570
157]	GTTCTTAAAGTAGTTTCAATTCCTAAGGAGACTTGGTATGATTTAGAAGAGGTT 1624
1625	5 CTGCTGGAAGAAATGACAGTTTTTCGGGAACCGACTGCTATTTCAGCAATGGAGCTTTCC 1684
1689	S ACTAAGCAGCAACAACTATATATTGGTTCAACGCTGCGGGTTCCCCAGCTCCCT 1738 ::: :::
1739	9 TTACACCGGTGTGATATTTACGGGAAAGCGTGTGCTGACTGTTGCCTCGCCCGAGACCCT 1798
1799	9 TACTGTGCTTGG 1810 9 TyrCysAlaTrpSerProProThrAlaThrCysValAlaLeuHisGlnThrGluSerPro 538
1811	1
4, 13	ACTGCAAAGAGACGCACAAGACGACAAGATAAGAAATGGAGACCCACTGACTCACTGT 19
0 9	TCAGACTTACACCATGATAATCACCATGCCCACAGCCCTGAAGAGAGAATCATCTATGGT 19
9 6	GTAGAGAATAGTAGCACATTTTGGAATGCGGAGAGTCGCAGAGAGGGCTGGTCTAT 20
2021	ATGAAGAG snGlyval
2081	1 ATCAGGACAGATCAAGGCCTTCTGCTAGGTAGCAAACAGAAGGATTCAGGCAATTAC 2140 :::
2141	1 CTCTGCCATGCGTGGAACATGGGTTCATACAAACTCTTCTTAAGGTA 2188

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GATCCTAAAAATCCAGTTGTATATGGAGTGTTTACGACTTCC-----AGTAACATTTTC 1204
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----IleIleGluGluLeuGlnIlePheProGlnGlyGlnProValGlnAsnLeuLeu 448
                                                                                                                                                                 ThrileValSerArgValAlaArgValCysLysGlyAspGluGlyGlyGluArgValLeu 242
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                 CGAACTCTTGGGCACCACCACCAATCAGGACAGAGCAGCATGATTCCAGGTGGCTCAAT 865
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                                                                                                                                                                                          Sequence 4, Application US/08556422A

Patent No. 657654

GENERAL INFORMATION:
APPLICANT: HALL, Kathryn T.
APPLICANT: SCHUITZE, Joachim L.
APPLICANT: SCHUITZE, Joachim L.
APPLICANT: BOUSSIOTIS, Vassiliki
APPLICANT: NUCLEIC ACIDS ENCODING CD100 MOLECULES
ITILE OF INVENTION: NUCLEIC ACIDS ENCODING CD100 MOLECULES
FILE REPERRENCE: DR.-005CPA2
CURRENT APPLICATION NUMBER: US/08/556,422A
CURRENT FILING DATE: 1999-11-09
NUMBER: OF SEQ ID NOS: 7
SOFTWARE: FastSEQ for Windows Version 4.0
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Mismatches:
Indels:
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Qy 1850 AGACGCACAAGACGACAAGATATAAGAAATGGAGACCCACTGACTCACTGTTCAGACTTA 1909	Db 508 SerArgProTrpThrGlnAsp1leGluGlyAlaSerValLysGluLeuCysLysAsn 526	Qy 1910 CACCATGATAATCACCATGGCCACAGGCCTGAAGAGAATCATCTATGGTGTAGAG 1966	Db 527SerSerTyrLysAlaArgPheLeuValProGlyLysPro 539.	Cy 1967	540 CysLysGlnValGlnIleGlnPr	2003	560 AsnLeualaintArgleunipvalhisasholyalakisovalasholarasholarasoc.vys	Addication in the contract of	Qy 2123 AAGGATTCAGGCAATTACCTCTGCGTGGAACATGGG 2164	Db 594 GlnGlyLeuGlyValPheGlnCysTrpSerIleGluGluGly 607	RESULT 12	7. Sequence 8, Application US/09653274	FALLICANT: BOYLE, Bryan J	; APPLICANT: Yeung, George Y ; APPLICANT: Arterburn, Matthew C	; APPLICANT: Mize, Nancy K ; APPLICANT: Tang, Y. Tom	, AFFLICANT: Dranac, Radoje T ; APPLICANT: Dranac, Radoje T ; TITLE OF INVENTION: Methods and Maaterials Relating to Semaphorin-Like	; TITLE OF INVENTION: Polypeptides and Polynucleotides ; FILE REFERENCE: HYS-23	; CURRENT APPLICATION NUMBER: US/09/653,274 ; CURRENT FILING DATE: 2000-08-31	FAILOR AFFILLATION NOTICE. ATTACKS OF OF IT MOST. 13	άi	42.	Scores:	2.73e-72 Length: 817.00 Matches: 47.54% Conservative:	st Local Similarity: 30.70% ery Match:	4 Gaps:	US-09-774-490-1 (1-2709) x US-09-653-274-8 (1-10/0) Qy 353 GCCAACAGCTCCAGTTATCATACCTTCCTTTTGGATGAGGAACGGAGTAGG 403		ricgacciggitaataicaaggaitti ::: :::	LeutyrilealaGlyArgAspGlnValTyrThrValAsnLeuAsnGluMet	ব'	6 tudoum

i TENGTH: 1086 i TYPE: PRT i CRGANISM: Homo sapiens US-09-653-274-4 Alignment Scores: 2.75e-72 Length: 1086 Score: 817.00 Matches: 206 Percent Similarity: 47.54% Conservative: 113 Best Local Similarity: 30.70% Mismatches: 233 Query Match: 46.69% Indels: 119 US-09-774-490-1 (1-2709) x US-09-653-274-4 (1-1086)	Qy 353 GCCAACAGCTCCAGTTATCATACCTTCCTTTTGGATGAGGAACGGAGTAGG 403 :::	Qy 464	Oy 560 CTTAAGGCATATAATCAGACTCACTTGTACGCCTGTGGAAGGGGGCTTTTCATCCAATT 619 125 PheValProArgAsnAspGluMetValPheValCySGlyThrAsnAlaPheAsnProMet 144 Oy 620 TGCACCTACATGAAATTGGACATCATCCTGAGGACAATTTTTTAAGCTGGAGAACTCA 679		Oy 728 ACAGCATCCTTTAATAGATGAGAATTATACTCTGGAACTGCAGCTGATTTTATGGGG 787 114 ASNVALALALEUDHAALAASDGIJUSEUTYTSERALATHTVALALAASDPHELEUALA 193 OY 788 CGAGACTTGCTATCTTGCACCCCCCCAATCAGGACAGCAT 847 DD 194 SerAspAlaValileTyTARGSERMetGIJASSGIYSERALALGHUSTYT 213	Oy 848 GATTCCAGGGGCTCAATGATCCAAAGTTCATTAGTGCCCACCTCATCTCAGAGAGTGC 907 214 AspSerLysTrplleLysGluProHisPheLeuHisAlalleGluTyrGly 230 Oy 908 AATCCTGAAGATGACAAAGTATTTTCTTCCGTGAAATGGATGATGAGAACAC :::
	Qy 1757 TACGGAAAGCGTGTGCTGACTCGCCCGAGACCCTTACTGTGCTTGGGAT 1813	Oy 1862Thraise Seral adlugly Tyrgluglina by Throluphed I	GTAGAGATAGTAGCACA 	Db 601 ProGlulleThrProLysval-IleAspThrTrpArgProLysLe 615 Qy 2042 GAAGAGGAAAGAAGATCAGAGTGGATGATCATATCATCAGGACAGATCAAGGCCTT 2101 Christofic in	Qy 2102 CTGCTACGTAGTC	RESULT 13 US-09-653-274-4 JUS-09-653-274-4 JUS-09-653-274-4 JUS-09-653-274-4 JUS-09-653-274-2 JUS-09-653-274-2 JUS-08-274-2 JUS-08-274-2 JUS-08-274-2 JUS-08-27-2 JUS-08-27-2 JUS-08-27-2 JUS-08-28-28-28-28-28-28-28-28-28-28-28-28-28

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ORGANISM: Homo sapiens
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                                       1196 AACATTTTCAAGGGATCAGCCGTGTGTATGTATAGCATGAGTGATGTGAGAAGGGTGTTC 1255
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                                                          380 LysThrSerlleAspPheProAspGluThrLeuSerPheIleLysSerHisProLeuMet 399
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APPLICANT: Mize, Nancy K
APPLICANT: Tang, Y. Tom
APPLICANT: Tang, Y. Tom
APPLICANT: Liu, Chenghua
APPLICANT: Liu, Chenghua
APPLICANT: Liu, Chenghua
APPLICANT: Division Methods and Maaterials Relating to Semaphorin-Like
TITLE OF INVENTION: Polypeptides and Polymucleotides
TITLE OF INVENTION: Polypeptides and Polymucleotides
TILE REFERENCE: HYS-23
CURRENT APPLICATION NUMBER: US/09/653,274
CURRENT APPLICATION NUMBER: 09/491,404
PRIOR FILING DATE: 2000-01-10
NUMBER OF SEQ ID NOS: 13
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 13
LENGTH: 641
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2138 TACCTCTGCCATGCGGTGGAACATGGGTTCA 2168
                                 655 eProLysGlyValArgTrpGluValGlnSer 665
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                                                                                                                                                           Sequence 13, Application US/09653274; Patent No. 6635742; GENERAL INFORMATION: APPLICANT: Boyle, Bryan J. APPLICANT: Yeung, George Y.
                                                                                                                                                                                                                                                   Boyle, Bryan J
Yeung, George Y
Arterburn, Matthew C
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US-09-774-490-1 (1-2709) x US-08-121-713D-62 (1-724)
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Best Local Similarity:
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               GATTCCAGGTGGCTCAATGATCCAAAGTTCATTAGTGCCCACCTCATCTCAGAGAGTGAC 907
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198 AspSerLysTrpIleLysGluProHisPheLeu-----HisAlaileGluTyrGly--- 214
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                                        GlnGlySerCysGlyArgValThrProGlyMetLeuLeuLeuThrGluAspPhePheAla 541
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                                                                                                                                     542 PheHisAsnHisSerAlaGluGlyTyrGluGlnAspThrGluPheGlyAsn---ThrAla
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/121,713D
FILING DATE: 13-SEP-1993
CLASSIFICATION: 514
ATTONINY AGENT INFORMATION:
NAME: OSman, Fichard A.
REGISTRATION NUMBER: 36,627
REFERENCE/DOCKET NUMBER: B94-002-1
TELEPROMUNICATION INFORMATION:
TELEPROMUNICATION INFORMATION:
TELEPROMUNICATION INFORMATION:
TELEPRAS: (415) 343-4342
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APPLICANT: Goodman, Alex L.
APPLICANT: Matthes, David
APPLICANT: Bentley, David R.
APPLICANT: Bentley, David R.
APPLICANT: O'Connor, Timothy
TTRE OF INVENTION: The Semaphorin Gene Family
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Mismatches:
Indels:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length:
Matches:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 62, Application US/08121713D Patent No. 5639856 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ZIP: 94104
COMPUTER REALABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                1955 TATGGT------
                                                                                                                                                                                                                                                                                                                                                                                1982 TIGGAAIGCAGICCGAAG 1999
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601 ProGluIleThrProLys 606
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LENGTH: 724 amino acids
TYPE: amino acid
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1370	aagaagtcatccago 	
1430	AATCCAGTGTTTCCTATGAACAATCGCCCAATAGTGATCAAAACGGATGTAAATTATCAA 1489 	
1490	TTTACACAAATHGTCGTAGACCGAGTGGATGCAGAAGATGGACAGTATGATGTT 1543	
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1658	ACTGCTATTTCAGCAATGGAGCTTTCCACTAAGCAGCAACAACTATATATTGGTTCAACG 1717	
1718	GCTGGGGTTGCCCAGCTCCCTTTACACCGGTGTGTGTTTACGGGAAAGCGTGTGCTGAG 1777 ::: ::	
1778	IGTIGCCTCGCCCGAGACCCTIACTGTGCTTGGGATGGTTCTGCATGTTCTCGC 1831	
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2054	GAAGAGATCAGAGTGGATGATCATATCATCAGGACAGATCAAGGCCTTCTGCTA 2107	
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September 23, 2004, 01:19:05; Search time 1239 Seconds (without alignments) 11035.942 Million cell updates/sec
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GenCore version 5.1.6
Copyright (c) 1993 - 2004 Compugen Ltd.
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

	-		Description	Sequence 1, Appli	Sequence 83, Appl	Sequence 283, App	Sequence 9, Appli				,	Sequence 347, App		Sequence 347, App	Sequence 347, App	347,	Sequence 347, App
SUMMARIES			ជ	US-09-774-490-1	US-10-247-671-83	US-10-097-340-283	US-10-262-538-9	US-10-067-632-53	US-10-404-438-2	US-09-946-374-309	US-10-015-395A-309	US-10-206-915-347	US-10-199-670-347	US-10-201-858-347	US-10-205-890-347	US-10-208-024-347	US-10-201-853-347
			DB	9	15	15	15	15	16	10	12	13	13	13	13	13	13
			Match Length DB	2709	2848	2530	2530	2601	2331	3871	3871	3871	3871	3871	3871	3871	3871
	οķο	Query	Match	100.0	92.9	95.6	92.6	95.6	26.5	26.5	26.5	26.5	26.5	26.5	26.5	26.5	26.5
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249 TECATICACCIGGITAATAICAAGGAITITCAAAAGAITGIGIGGCCAGTAICTIACAC 308 493 CAGAAGAGAIGAATGGAAGAGGTGGAAAAGAATGCCIGAAAAAAATGTGTGGCTAAITCAI 552	CAGAAGAATGAATGAATGCAAGTGGGCTGGAAAAGACATCCTGAAGAATGTCCTAATTTCAT 36	553 CARGIACTTAAGGCATATVAATCAGACTCACTTGTACGCCTGTGGAACGGGGGGCTTTTCA 612	613 TCCAATTTGCACCTACATTGAAATTGGACATCATCCTGAGGACAATATTTTAAGCTGGA 672 	673 GAACTCACATTTTGAAAACGGCCGTGGGAAGAGTCCATATGACCCTAAGCTGCTGACAGC 732 	733 ATCCCTITTAATAGATGGAGAATTATACTCTGGAACTGCAGCTGATTTATGGGGCGAGA 792 	793 CTTTGCTATCTTCCGAACTCTTGGGCACCACCAATCAGGACAGAGAGCAGCATGATTC 852 	853 CAGGTGGCTCAATGATCCAAAGTTCATTAGTGCCCACCTCATCTCAGAGAGTGACAATCC 912	913 TGAAGATGACAAAGTATACTTTTTTTCTTCCGTGAAATGCAATAGATGGAGAACACTCTGG 972 	973 AAAAGCTACTCACGCTAGAATAGGTCAGATATGCAAGAATGACTTTGGAGGGCACAGAAG 1032 	1033 TCTGGTGAATAAATGGACAACATTCCTCAAAGGTCGTCTGATTTGCTCAGTGCCAGGTCC 1092	1093 AAATGGCAITGACACICAITITGAIGAACTGCAGGAIGIAITCCTAAIGAACTITAAAGA 1152 	1153 TCCTAAAAATCCAGTTGTATATGGAGTGTTTACGACTTCCAGTAACATTTCAAGGGATC 1212 	1213 AGCCGTGTGTATGTGTGAGTGATGTGAGAGGGTGTTCCTTGGTCCATATGCCCA 1272	1273 CAGGGATGGACCCAACTATCAATGGGTGCCTTATCAAGGAAGAGTCCCCTATCCACGGCC 1332 	1333 AGGAACTTGTCCCAGCAAACATTTGGTGGTTTTGACTCTACAAAGGACCTTCCTGATGA 1392	1393 IGTIATAACCTTIGCAAGAAGICAICCAGCCAIGTACCAATCCAGIGTITCCTAIGAACAA 1452 	1453 TGGCCCAATAGTGATCAAACGGATGTAAATTATCAATTTACACAAATTGTCGTAGACCG 1512	
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93 GENERAL CONTROLLES
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Db 2409 GAACTTTTTAAAN RESULT 4 (B-10-262-538-9) SECONDITALAN (B-10-262-538-9) SECONDITAL) SUBLICATION NO. US200301) GENERAL INFORMATION: NEU FILE PEPERENCE: 28967/3 CURRENT FILING DATE: 2 UNDHER OF SEQ ID NOS: 3 SOFTWARE: PATENTION NUMBER OF SEQ ID NOS: 3 SOFTWARE: PATENTION NOW CURRENT FILING DATE: 2 UNDHER OF SEQ ID NOS: 3 SOFTWARE: PATENTION NOW PEATURE: NAME/ CURRENT FILING DATE: 2 UNDHER OF SEQ ID NOS: 3 SOFTWARE: PATENTION NOS: 3 SOFTWARE: PATENTION NOS: 3 CURRENT FILING DATE: 2 UNDHER OF SEQ ID NOS: 3 SOFTWARE: PATENTION NOS: 3 SOFTWARE: PATENTION NOS: 3 CURRENT FILING DATE: 2 UNDHER OF SEQ ID NOS: 3 SOFTWARE: PATENTION NOS: 3 CURRENT PATENTION OUR DATE DO AGAGGCAACTAC ON 373 TACCTTCCTTTT ON 373 TACCTTCCTTTT ON 493 CAGAAGAATGA ON 373 TACCTTCCTTTTAAT ON 499 GAACTCACATTTAAT ON 733 ATCCTTTTAAT ON 733 ATCCTTTTTAAT ON 733 ATCCTTTTTAAT ON 733 ATCCTTTTTAAT ON 733 ATCCTTTTAAT ON 733 ATCCTTTTTAAT ON 733 ATCCTTTTTTAAT ON 733 ATCCTTTTTAAT ON 734 ATCCTTTTTAAT ON 734 ATCCTTTTTAAT ON 734 ATCCTTTTTAAT ON 744 O

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COMPUTER: IMPRO COMPUTER: I
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92.6%; Score 2508.4;
Best Local Similarity 100.0%; Pred. No. 0;
Matches 2509; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LOCATION: 16..2331
SEQUENCE DESCRIPTION: SEQ ID NO: 53;
                                       ZIP: 94104
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEPHONE: (415)343-4341
TELEFAX: (415) 343-4342
TELEX: «DIKNOM»>
INFORMATION FOR SEQ ID NO: 53: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 2601 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
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LOCATION: 16..
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                                                                                                                     1749 CAGCCCTGAAGAAGAATCATCTATGGTGTAGAGAATAGTAGCACATTTTTGGAATGCAG
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US-10-067-632-53

US-10-067-632-53

Sequence 53, Application US/10067632

Publication No. US20030166849A1

GENERAL INFORMATION:

KOLOGKIN, Alex L.

Matches, David

Bentley, David

COCORDO, THE OF INVENTION:

NUMBER OF SEQUENCES: 100

CORRESPONDENCE ADDRESS:

ADDRESSEE: SCIENCE & TECHNOLOGY LAW GROUP

STREET: 268 Bush Street, Suite 3200

CITY: San Francisco
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	733	atcctttpatagatggagattatactctggaactgcagctgatttatggggcgaga 792 	
	793 (CITIGCIARCITCCGAACTCTTGGGCACCACCAATCAGGACAGAGGAGAGATC 852 	
	853	CAGGTGGCTCAATGATCCAAAGTTCATTAGTGCCCACCTCATCTCAGAGAGTGACAATCC 912 	
	913	TGAAGATGACAAAGTATACTTTTCTTCCGTGAAAATGCAATAGATGGAGAACACTCTGG 972 	
	973 7	AAAAGCTACTCACGCTAGAATAGGTCAGATATGCAAGAATGACTTTGGAGGGCACAGAAG 1032 	
.7	849	TCTGGTGAATAAATGGACAACATTCCTCAAAGCTCGTCTGATTTGCTCAGTGCCAGGTCC 1092 	
, 3	1093	AAATGGCATTGACACTCATTTTGATGAACTGCAGGATGTATTCCTAATGAACTTTAAAGA 1152 	
.,	969	TCCTAAAATCCAGTIGIAIAIGGAGIGITIACGACTTCCAGTAACATITTCAAGGGAIC 1212 	
	1213,7	AGCCGTGTGTATGTATAGCATGAGTGTGAGAAGGGTGTTCCTTGGTCCATATGCCCA 1272.	
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ન ન	333	aggaacttgtcccagcaaacatttggtggtttgactcracaaaggaccttcctgatga 1392 	
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н н	453	TCGCCCAATAGTGAICAAAAACGGATGTAAATTATCAATTTACACAAATTGTCGTAGACG 1512 	
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1569 TATTTACGGGAAAGCGTCTGCTGAGTCTTGCCTCGCCCGAGACCTTACTGTGCTTGGGA 1628
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US-10-404-438-2

Sequence 2, Application US/10404438

Publication to US20030232411A1

GENERAL INFORMATION:

APPLICANT: Ono Pharmaceutical Co., Ltd.

APPLICANT: Ono Pharmaceutical Co., Ltd.

TITLE OF INVENTION: NOVEL POLYPEPTIDE, cDNA ENCODING THE SAME, AND USE THEREOF

FILE REPERENCE: Q59878

CURRENT APPLICATION NUMBER: US/10/404,438

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Pred. No. 5e-206;
0; Mismatches 739; Indels
CURRENT FILING DATE: 2003-04-02
PRIOR APPLICATION NUMBER: US/09/582,419A
PRIOR FILING DATE: 2000-06-26
PRIOR APPLICATION NUMBER: PCT/UP98/05952
PRIOR FILING DATE: 1998-12-25
PRIOR FILING DATE: 1997-12-26
NUMBER OF SEQ ID NOS: 28
SOFTWARE: Patentin version 3.1
SEQ ID NO 2
                                                                                                                                                                                                                                                                                                                                                                                    26.5%;
61.9%;
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Best Local Similarity 61.9
Matches 1233; Conservative
                                                                                                                                                                                                                                                                                 TYPE: DNA
ORGANISM: Homo sapiens
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PRIOR FILING DATE: 1998-09-10
PRIOR APPLICATION NUMBER: 60/099815
PRIOR PELING DATE: 1998-09-10
PRIOR PLING DATE: 1998-09-10
PRIOR PLING DATE: 1998-09-15
PRIOR PLING DATE: 1998-09-16
PRIOR PLING DATE: 1998-09-17
PRIOR APPLICATION NUMBER: 60/10019
PRIOR PLING DATE: 1998-09-17
PRIOR PLING DATE: 1998-09-17
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PRIOR PLING DATE: 1998-09-18
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PRIOR FILING DATE: 1998-09-24
PRIOR APPLICATION NUMBER: 60/102207
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FILING DATE: 1998-09-24
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CURRENT APPLICATION NUMBER: 05/098716

PRIOR APPLICATION NUMBER: 06/098716

PRIOR PILING DATE: 1938-09-01

PRIOR PILING DATE: 1938-09-01

PRIOR PILING DATE: 1938-09-01

PRIOR PILING DATE: 1938-09-01

PRIOR APPLICATION NUMBER: 06/09803

PRIOR APPLICATION NUMBER: 06/09803

PRIOR PILING DATE: 1938-09-01

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PRIOR PILING DATE: 1938-09-02

PRIOR PILING DATE: 1938-09-02

PRIOR PILING DATE: 1938-09-09

PRIOR PI
   2174 ACTOTICITAAGGIAACCCIGGAAGICATIGACACAGAGCATITGGAAGAACTICTICAT 2233
                                                                   2023 ACCATAGTGAAGCTGACTTTGAATGTCATTGAGAATGAACAGATGGAAAATACCCAGAGG 2082
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Fong, Sherman
God, Wal-Qiang
Goddard, Audrey
Godowski, Paul J.
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APPLICATION NUMBER: 60/099792
FILING DATE: 1998-09-10
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Roy, Margaret Ann
Smith, Victoria
Stewart, Timothy A.
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Williams, P. Mickey
Wood, William I.
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Hillan, Kenneth J
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Eaton, Dan L.
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9-29	<u>ਰ</u>	215 ACTITGAAGCAAAATATTCCAAGACTCAAGCTAACCTACAAAGACTTGCTGCTTTCAAAT 274
FILING DATE	Å,	329 AATGIGALCACTITCAATGGCTTGGCCAACAGCTCCAGTTATCATACCTTCCTT
	qa	275 ACCTGTATTCCCTTTTTGGGTCATCATCAAAGGACTGGATTTTCAAACTCTTCTCTTGGAT 334
	ò	389 GAGGAACGGAGTAGGCTGTATGTTGGAGGAAAGGATCACATATTTTCATTCGACCTGGTT 448
	qo	335 GAGGAAAGGAGGCAGGCTCTTGGGAGCCAAAGACCACATTCTACTCAGTCTGGTT 394
	ζ	449 AATATCAAGGATTTTCAAAAGATTGTGTGGCCAGTATCTTACACCAGAAGAGATGA 505
	업	395 GACTTRAACAARAATTTTRAGAAGATTTRTTGGCCTGCRAAGGAACGGGTGGAATTA 454
PRIOR FILING DATE: 1998-09-30 PRIOR APPLICATION NUMBER: 60/102571	ò	506 TECAAGTEGGGCTEGAAAAGACATCCTGAAAGAATGTGCTAATTTCATCAAGGTACTTAAG 565
	qa	455 IGTABATTAGCTGGGAAAGATGCCAATACAGAATGTGCAAATTTCATCAGAGTACTTCAG 514
PRIOR FILING DATE: 1998-10-01 PRIOR APPLICATION NUMBER: 60/102687	ò	566 GCATATAATCAGACTCACTTGTACGCCTGTGGAACGGGGGCTTTTCATCCAATTTGCACC 625
PRIOR FILING DATE: 1998-10-01 PRIOR APPLICATION NUMBER: 60/102965	qu	515 cécharakakatékakanarangnengiagakatagakéarirtéarékaharérese 574
PRIOR FILING DATE: 1998-10-02 PRIOR APPLICATION NUMBER: 60/103258	δò	TACATTGAAATTGGACATCATCCTGAGGACAATATTTTTAAGCTGGAGAACTCACATTTT
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PRIOR APPLICATION NUMBER: 60/103315	ò	
PRIOR FILING DATE: 1998-10-07 PRIOR APPLICATION NUMBER: 60/103328	qa	635 GAGTCTGGCAGACTGAAATGTCCTTTCGATCCTCAGCAGCTTTTGCTTCAGTAATGACA 694
PRIOR FILING DATE: 1998-10-07 PRIOR APPLICATION NUMBER: 60/103395	ò	746 GAIGGAGAAITAIACICIGGAACIGCAGCIGAITITAIGGGGGGAGACITICCIAICITC 805
FILING DATE: 1998-10 APPLICATION NUMBER:	qq	695 GAIGAGTACCTCTACTCTGGAACAGCTTCTGATTTTCCTTGGCAAGATACTGCATTCACT 754
PRIOR FILING DATE: 1998-10-07 PRIOR APPLICATION NUMBER: 60/103401	λō	806 CGAACTCTTGGGCACCACCACCCAATCAGGACAGAGCAGCATGATTCCAGG 856
	QO	755 GGATCCCTTGGGCCTACTCATGACCACTACATCAGAACTGACATTTCAGAGCACTAC 814
FILING DATE: 1998-10-0 APPLICATION NUMBER: 60	Λō	857 TGGCTCAATGATCCAAAGTTCATTAGTGCCCACCTCATCTCAGAGAGTGACAATCCTGAA 916
	Q O	815 TGGCTCAATGGAAGATTTATTGGAACTTTCTTCATACCAGACACCTACAATCCAGAT 874
FILING DATE: 1998-10-08 APPLICATION NUMBER: 60/10367	ò	917 GATGACAAAGTATACTITTCTICCGTGAAAAIGCAATAGATGGAGAACACTCTGGAAAA 976
	qa	875 GAIGATAAATATATTTCTTTTCTTGAATCATCTCAAGAAGGCAGTACCTCCGATAAA 934
	ò	977 GCTACTCACGTAGAATAGGTCAGATATGCAAGAATGACTTTGGAGGCACAGAAGTCTG 1036
	qq	935 ACCATCCTTTCTCGAGTTGGAAGAGTTTGTAAGAATGATGTAGGAGGACAACGCAGCCTG 994
PRIOR FILING DATE: 1998-10-07 PRIOR ADDITIONAL MANAGED. 60/10c000	ò	1037 GTGAATAAATGGACAACATTCCTCAAAGCTCGTCTGATTTGCTCAGTGCCAGGTCCAAAT 1096
FILING DATE: 1998-10	අු	995 ATAAACAAGTGGACGACTTTTCTTAAGGCCAGACTGATTTGCTCAATTCCTGGAAGTGAT 1054
PRIOR APPLICATION NUMBER: 60/105002 PRIOR FILING DATE: 1998-10-20	<u></u>	1097 のものは、それのようなななななななななななななななななななななななななななななななななななな
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	qq	1115 AGAAATCCTGTAGTATATGGAGTCTTTACTACAACCAGCTCCATCTTCAAAGGCTCTGCT 1174
	ζζ	1217 GIGIGIAIGIAIAGCAIGAGIGAIGAGAAGGAGGIGTICCTIGGICCAIAIGCCCACAGG 1276
PRIOR FILING DATE: 1998-10-26 PRIOR APPLICATION NUMBER: 60/105807	qq	1175 GTTTGTGTGTATAGCATGGCTGACATCAGAGCAGTTTTTAATGGTCCATATGGTCTCATAAG 1234
26.5%;	λσ	1277 GAIGGACCCAACTAICAAIGGGIGCCITAICAAGGAAGAGICCCCTAICCACGGCCAGGA 1336
Pred. No. 7.4e-206; Tagal 21. Care	qa	1235 GAAAGTGCAGACCATGGTTGGGGGGTGTAGGTGGGAGAATTCCTTATCCACGGCCTGGT 1294
vative U; Mismatch		

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TITLE OF INVENTION: Acids Encoding the Same
FILE REFERENCE: P2830P1C57
CURRENT APPLICATION NUMBER: US/10/015, 395A
CURRENT APPLICATION FORDE: 2001-12-12
Prior application removed - See file Wrapper or Palm
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Gao, Wei-Qiang
Goddard, Audrey
Godowski, Paul J.
Grimaldi, Christopher J.
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Best Local Similarity 61.9<sup>5</sup>
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APPLICANT: Botstein, David
APPLICANT: Desnoyers, Luc
APPLICANT: Eaton, Dan I.
APPLICANT: Ferrara, Napoleone
APPLICANT: Fong, Sherman
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2234 AAAGATGATGATG 2246
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GCTACTCACGCTAGAATAGGTCAGATATGCAAGAATGACTTTGGAGGGGCACAGAAGTCTG 103 ACATCCTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTT	TGGCTCAATGGAGCAAATTTATTGGAACTTTCTTCATACCAGACACCTACAATCCAGAT 87 GATGACAAAGTATACTTTCTTCCGTGAAAATGCAATAGATGGAAAAGAAGAACACTTGGAAAA 87 [

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1355 GTCATCAGTTTCATAAAGCGGCACTCTGTGATGTATAAGTCCGTATACCCAGTTGCAGGA 1414
                                                                      1454 CGCCCCAATAGTGATCAAAACGGATGTAAATTATCAATTTACACAAATTGTCGTAGACCGA 1513
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RESULT 10

US-10-199-670-347

Sequence 347, Application US/10199670

Publication No. US20040033560A1

APPLICANT: Baker, Kevin P.

APPLICANT: Chen, Jian

APPLICANT: Goddard, Audrey

APPLICANT: Goddard, Audrey

APPLICANT: Goddard, Audrey

APPLICANT: Goddard, Audrey

APPLICANT: Goddwski, Paul J.

APPLICANT: Goddwski, Paul J.

APPLICANT: Smith, Victoria

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977 GCTACTCACGCTAGAATAGGTCAGATATGCAAGAATGACTTTGGAGGGCACAGAAGTCTG 1036
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                TGGCTCAATGGAGCAAAATTTATTGGAACTTTCTTCATACCAGACACCTACAATCCAGAT
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                                                                                AND NUCLEIC
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APPLICANT: Watanabe, Colin K.
APPLICANT: Wood, William I.
APPLICANT: ADAG, Zemin I.
ITILE OF INVENTION: ECRETED AND TEANSWENBRANE POLYPEPTIDES AND ITILE OF INVENTION: ACIDS ENCODING THE SAME
TITLE OF INVENTION: ACIDS ENCODING THE SAME
ITILE OF INVENTION: ACIDS ENCODING THE SAME
CURRENT FILING DATE: 12002-07-19
FRICH PAPLICATION NUMBER: 10/052586
PRIOR FILING DATE: 1997-09-18
PRIOR APPLICATION NUMBER: 60/059263
PRIOR APPLICATION NUMBER: 60/059266
PRIOR APPLICATION NUMBER: 60/063250
PRIOR PILING DATE: 1997-09-18
PRIOR PILING DATE: 1997-10-24
PRIOR PILING DATE: 1997-10-24
PRIOR APPLICATION NUMBER: 60/063120
PRIOR APPLICATION NUMBER: 60/063121
PRIOR APPLICATION NUMBER: 60/063121
PRIOR APPLICATION NUMBER: 60/063540
PRIOR PILING DATE: 1997-10-24
PRIOR PILING DATE: 1997-10-28
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 718.6; DB 13;
Pred. No. 7.4e-206;
0; Mismatches 739;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity 61.9%;
Matches 1233; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; TYPE: DNA
; ORGANISM: Homo Sapien
US-10-199-670-347
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NUMBER OF SEQ ID NOS: 612
                        ; SEQ ID NO 347
; LENGTH: 3871
; TYPE: DNA
; ORGANISM: HOMO Sapien
US-10-201-858-347
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                                                1874 AGAAATGGAGCCCACTGACTCACTGTTCAGACTTACACCATGATAATCACCATGGCCAC 1933
                                                                                                1832 AAATATGGCGACCCAATCACCCCAGTGCTGGGACATCGAAGACAGCATTAGTCATG---AA 1888
                                                                                                                                                   1934 AGCCCTGAAGAGAATCATCTATGGTGTGAGAATAGTAGCACATTTTTGGAATGCAGT 1993
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                                                                                                                                                                                                                                                                                               1949 CCTAAATCCCAACAAGCAACTATTAAATGGTATATCCAGAGGTCAGGGGGATGAGCATCGA 2008
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TITLE OF INVENTION: ACIDS ENCEPTED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: P3430812428
CURRENT APPLICATION NUMBER: US/10/201, 658
PRIOR FILING DATE: 2002-07-23
PRIOR FILING DATE: 10/05266
PRIOR FILING DATE: 10/05-08-18
PRIOR PLILING DATE: 1997-09-18
PRIOR PLILING DATE: 1997-09-18
PRIOR PLILING DATE: 1997-10-17
PRIOR PLILING DATE: 1997-10-17
PRIOR PLILING DATE: 1997-10-24
PRIOR PLILING DATE: 1997-10-28
1772 GGAAATGCATGCTCTCGATATGCTCCTACTTCTAAAAGGAGAGCTAGACGCCAAGATGTA 1831
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FILING DATE: 1997-10-28
Application data removed - See File Wrapper or PALM.
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Publication No. US2004038337A1
GENERAL INFORMATION:
APPLICANT: Baker Kevin P.
APPLICANT: Chen, Jian
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Watanabe, Colin K.
Wood, William I.
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Godowski, Paul J.
Gurney, Austin L.
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APPLICANT:
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1037 GIGAATAAAIGGACAACATICCICAAAGCICGICIGAITIGCICAGGIGCCAGGICCAAAI 1096
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                                                                                                                                                                                   269 AATGGGAAGAACAATGTGCCAAGGCTGAAATTATCCTACAAAGAAATGTTGGAATCCAAC 328
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                                                                                                                                                                                                                                                                 215 actiticaagcaaaataticcaagacticaagciaacctacaaagactigcigcuticaaar 274
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             686 GAAAACGGCCGTGGGAAGAGTCCATATGACCCTAAGCTGCTGACAGCATCCCTTTTAATA
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                                                                                            21; Gaps
         Length 3871;
Score 718.6; DB 13; Length Pred. No. 7.4e-206; 0; Mismatches 739; Indels
Query Match
Best Local Similarity 61.9%;
Matches 1233; Conservative (
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1115 AGAAATCCTGTAGTATATGGAGTCTTTACTACAACCAGCTCCATCTTCAAAGGCTCTGCT 1174

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TYPE: DNA
ORGANISM: Homo Sapien
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 GTGTGTATGTATAGCATGAGTGATGTGAGAAGGGTGTTCCTTGGTCCATATGCCCACAGG 1276
                                                                      GATGGACCCAACTATCAATGGGTGCCTTATCAAGGAAGAGTCCCCTATCCACGGCCAGGA 1336
                                                                                                            GADAGTGCAGCCATCGTTGGGTGCAGTATGATGGGAGAATTCCTTATCCACGGCCTGGT 1294
                                                                                                                                                   ACTIGICCCAGCAAAACATTIG----GIGGITTIGACICTACAAAGGACCTICCIGAIGAT 1393
                                                                                                                                                                                    ACATGICCAAGCAAAACCIATGACCCACIGATIAAGICCACCGAGAITTTICCAGATGAT 1354
                                                                                                                                                                                                                          GTTATAACCITIGCAAGAAGICAICCAGCCAIGIACAAICCAGIGIITCCIAIGAACAAI 1453
                                                                                                                                                                                                                                                             GICAICAGITICATAAAGCGGCACTCTGTGATGTATAAGTCCGTATACCCAGTTGCAGGA 1414
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                                     GTTTGTGTGTATAGCATGGCTGACATCAGAGCAGTTTTTAATGGTCCATATGCTCATAAG
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GENERAL INFORMATION:

APPLICANT: Baker, Kevin P.

APPLICANT: Chen, Jian P.

APPLICANT: Chen, Jian P.

APPLICANT: Goddard, Audrey

APPLICANT: Goddard, Audrey

APPLICANT: Goddard, Audrey

APPLICANT: Goddard, Laul J.

APPLICANT: Goddard, Laul J.

APPLICANT: Word, William I.

APPLICANT: Word, William I.

APPLICANT: Word, William I.

APPLICANT: Word, William I.

APPLICANT: Alang, Zemin

TITLE OF INVENTION: SECRETED AND TRANSKEMBRANE POLYPEPTIDES AND NUCLEIC

TITLE DEPRENOR: P3430R1C519

APPLICANT: ALANG ZOND: 105,266

PRIOR PLILNG DATE: 2002-01-15

PRIOR PLILNG DATE: 1997-00-18

PRIOR APPLICATION NUMBER: 60/063266

PRIOR PLILNG DATE: 1997-10-24

PRIOR PLILNG DATE: 1997-10-28

PR 269 AATGGGAAGAACAATGTGCCCAAGGCTGAAATTATCCTACAAAGAAATGTTGGAATCCAAC 328 GAGGAAAGAGGCAGCCTCTTGGGAGCCAAAGACCACATCTTTCTACTCAGTT 394 565 215 ACTTTGAAGCAAAATATTCCAAGACTCAAGCTAACCTACAAAGACTTGCTGCTTTCAAAT 274 AGCTGTATTCCCTTTTTGGGTTCATCAGAAGGACTGGATTTTCAAACTCTTCTTAGAT 334 449 AA---TATCAAGGATTTTCAAAAGATTGTGTGGCCAGTATCTTACACCAGAAGAGATGAA 505 TGCAAGTGGGCTGGAAAAGAATCCTGAAAGAATGTGCTAATTTCATCAAGGTACTTAAG GAGGAACGGAGTAGGTTGTTGGAGCAAAGGATCACATATTTTCATTCGACCTGGTT 395 GACTTAAACAAAATTTTAAGAAGATTTATTGGCCTGCTGCAAAGGAACGGGGGATTA Gaps Length 3871; 21; Query Match 26.5%; Score 718.6; DB 13; Length Best Local Similarity 61.9%; Pred. No. 7.4e-206; Matches 1233; Conservative 0; Mismatches 739; Indels Application US/10205890 5. US20040048334A1

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                                                 1592 GAGTIGCAGAIATICAAGCACTCATCAATCÁTCÍTGAACÁTGGAATIGÍCTCTGAAGCAG
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RESULT 13
REPLICANT: Gene, Jian
APPLICANT: Gene, Jian
APPLICANT: Geddard, Audrey
APPLICANT: Geddowski, Paul J.
APPLICANT: Gene, Jian
APPLICANT: Gurney, Austin L.
APPLICANT: Matanabe. Colin K.
APPLICANT: Watanabe. Colin K.
APPLICANT: Matanabe. Colin K.
APPLICANT: Modd, William I.
TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
TITLE REFERENCE: P3430R1C38
CURRENT PAPLICATION NUMBER: 105/10286
PRIOR FILING DATE: 2002-01-15
PRIOR FILING DATE: 2002-01-15
PRIOR FILING DATE: 1997-09-18
PRIOR FILING DATE: 1997-09-18
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Pred. No. 7.4e-206;
0; Mismatches 739;
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PRIOR APPLICATION NUMBER: 60/062250
PRIOR FILING DATE: 1997-10-17
PRIOR PILING DATE: 1997-10-17
PRIOR PILING DATE: 1997-10-24
PRIOR PILING DATE: 1997-10-24
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PRIOR PILING DATE: 1997-10-24
PRIOR PILING DATE: 1997-10-24
PRIOR PILING DATE: 1997-10-28
PRIOR SPELICATION NUMBER: 60/063544
PRIOR PILING DATE: 1997-10-28
PRIOR SPELING DATE: 1997-10-28
PRIOR 347
PRIOR 347
                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity 61.9%;
Matches 1233; Conservative
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US-10-208-024-347
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APPLICANT: Watenabe_Colin K.
APPLICANT: Wood, William I.
APPLICANT: Wood, William I.
APPLICANT: Wood, William I.
APPLICANT: Zanag, Zenin
TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE SEPREBRENCE: P340R1C465
CURRENT APPLICATION NUMBER: US/10/2286
PRIOR APPLICATION NUMBER: US/05286
PRIOR APPLICATION NUMBER: 00/059263
PRIOR APPLICATION NUMBER: 60/059266
PRIOR FILING DATE: 1997-09-18
PRIOR FILING DATE: 1997-09-18
PRIOR FILING DATE: 1997-10-24
PRIOR APPLICATION NUMBER: 60/063120
PRIOR APPLICATION NUMBER: 60/063120
PRIOR FILING DATE: 1997-10-24
PRIOR FILING DATE: 1997-10-28
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                                                                                                                                                                    ACCATAGTGAAGCTGACTTTGAATGTCATTGAGAATGAACAGATGGAAAATACCCCAGAGG 2188
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NUMBER OF SEQ ID NOS: 612
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Pred, No. 7.4e-206;
0; Mismatches '739;
                                                                                                                                                                                                                                                                                                                                 Sequence 347, Application US/10201853 Publication No. US20040053358A1 GENERAL INFORMATION:
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Best Local Similarity 61.9%;
Matches 1233; Conservative
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Godowski, Paul J.
Gurney, Austin L.
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AND NUCLEIC
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                               APPLICANT: Watanabe, Colin K.
APPLICANT: Wood, William I.
APPLICANT: Zhang, Semin
TITLE OF INVENTION: SECRETED AND TRAITILE OF INVENTION: ACIDS ENCODING
FILE REFERENCE: P3430R1C41
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APPLICATION NUMBER: 60/080327
FILING DATE: 1998-04-01
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APPLICATION NUMBER: 60/080194
Smith, Victoria
Watanabe, Colin K.
Wood, William I.
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   GITATAACCITIGCAAGAAGICATCCAGCCATGIACAAICCAGIGITIGCTAIGAACAAI 1453
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                                                                           1355 Grearchentraraaaceecacacrererearenaraareeceraraeceea
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                                                                                                                                                      CGCCCAATAGTGATCAAAACGGATGTAAATTAATATTACACAAATTGTCGTAGACCGA
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US-10-174-581-347
Sequence 347, Application US/10174581
Sequence 347, Application US/10174581
GENERAL INFORMATION:
APPLICANT: BAREF, Kevin P.
APPLICANT: Chen, Jian
APPLICANT: Goddard, Audrey
APPLICANT: Goddard, Audrey
APPLICANT: Godowski, Paul J.
APPLICANT: Gunney, Austin L.
APPLICANT: Pan, James
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2234 AAAGATGATG 2246
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PRIOR APPLICATION NUMBER: 60/080333
PRIOR FILING DATE: 1998-04-01
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PRIOR PELING DATE: 1998-04-08
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PRIOR PELING DATE: 1998-04-15
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PRIOR PELING DATE: 1998-04-21
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PRIOR PELING DATE: 1998-05-07
PRIOR PELING DATE: 1998-05-15
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FILING DATE: 1998-06-04
APPLICATION NUMBER: 60/088167
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PRIOR FILING DATE: 1998-06-05
PRIOR APPLICATION NUMBER: 60/086217
PRIOR PLING DATE: 1998-06-05
PRIOR PLING DATE: 1998-06-05
PRIOR PAPLICATION NUMBER: 60/08652
PRIOR PAPLICATION NUMBER: 60/08655
PRIOR PILING DATE: 1998-06-09
PRIOR PILING DATE: 1998-06-09
PRIOR PLING DATE: 1998-06-10
PRIOR APPLICATION NUMBER: 60/08673
PRIOR PLING DATE: 1998-06-10
PRIOR PELING DATE: 1998-06-10
PRIOR PLING DATE: 1998-06-11
PRIOR PLING DATE: 1998-06-12
PRIOR PLING DATE: 1998-06-12 PRIOR FILING DATE: 1998-06-16 PRIOR APPLICATION NUMBER: 60/089514 PRIOR FILING DATE: 1998-06-16 APPLICATION NUMBER: 60/089598 FILING DATE: 1998-06-17 APPLICATION NUMBER: 60/089653 LICATION NUMBER: 60/088202 ING DATE: 1998-06-05 APPLICATION NUMBER: 60/089538 998-06-17

21; DB 13; Length 3871; Indels 7.4e-206; nes 739; Query Match 26.5%; Score 718.6; Best Local Similarity 61.9%; Pred. No. 7.4e Matches 1233; Conservative 0; Mismatches

388 274 334 448 394 454 269 AATGGGAAGAACAATGTGCCAAGGCTGAAATTATCCTACAAAGAAATGTTGGAATCCAAC 215 ACTTTGAAGCAAAATATTCCAAGACTCAAGGCTAACCTACAAAGACTTGCTGCTTTCAAAT 395 GACTTAAACAAAATTTTAAGAAGATTTATTGGCCTGCAAAGGAACGGGGGGGAATTA 275 AGCTGTATTCCCTTTTTGGGTTCATCAGAGGACTGGATTTTCAAACTCTTCTTAGAT 389 GAGGAACGGAGTAGGTGTTGGAGCAAAGGATCACATATTTTCATTCGACCTGGTT 335. GAGGAAAGAGGCAGCTGCTTGGGAGCCAAAGACGACATCTTTCTACTCAGTCTGGTT 449 AA---TATCAAGGATTTTCAAAAGATTGTGGGCCAGTATCTTACACCAGAAGAGATGAA ò a 8 a qq ò g

TGCAAGTGGGCTGGAAAAGACATCCTGAAAGAATGTGCTAATTTCATCAAGGTACTTAAG 206 q

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S66 GCATATAATCAGACTCACTTGTACGCCTGTGGAACGGGGCTTTTCATCCAATTTGCACC 625 515 cécrarakacakakérékenharkardrekékakérékakaritrékrécakhardrege 574 455 IGTAAATTAGCTGGGAAAGATGCCAATACAGAATGTGCAAATTTCATCAGAGTACTTCAG

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Db 1652 CAACAATTG Qy 1754 ATTTACGGG Qy 1712 ACTTATGGG Qy 1814 GGTTCTGCA Qy 1872 GGAATGCA Qy 1874 AGAATGGG	Oy 1934 AGCCCTGAA Db 1889 ACTGCTGAT QY 1994 CCGAAGTCG QY 2054 GAAGAGATC QY 2054 GAAGAGATC QY 2059 GAAGAGATC QY 2114 CTACAACAG QY 2114 CTACACAGAG QY 2114 ACTCTTCTT Db 2059 ACCATAGTG	2189 ; arch complet b time : 126	· ·
626 TACATTGAAATTGGACATCTCTGAGGACAATTTTTAAGCTGGACAACTCACATTTT 685	TGGCTCAATGGAGCAAATTTATTGGAACTTTCTTCATACAGAGAGTGACAATCCTGAATGGACTCAATGGAGAGTGAATCTTTCTT	1157 AANAATCCAGTTGTATATGGAGTTTTACGACTTCCAGTAACATTTTCAAGGGATCAGCC 1216 1115 AGAAATCCAGTTGTATATGGAGTGTTTACTACAGTGCTTCTTCTTCTTCTAAGGGATCAGCC 1216 1115 GTGTGTATGTTAGGAGTGGTTGTAGGAGAGGGTGTTCTTCGAAGGCTCTGCT 1174 1217 GTGTGTATGTTAGCATGGAGAGGGTGTTCTTGGTCCATATGCCCAGGG 1276 1175 GTTTGTGTGTATAGCATGGAGAGAGGGTGTTTTAATGGTCCATATGCCCAGGG 1234 1277 GATGGACCCAACTATGGAGAGAGTGTTTTAATGGTCCATATGCTCATAGG 1234 1285 GAAAGTGCAACAACTATGGAGAGAGAGAGTTTTTAATGGTCCATATGCTCATAGG 1294 1387 ACTTGTCCCAGCAAACATTTGAGTGGTTTTGACTCTAATGCAGGCCTGGT 1393 1295 ACATGTCCCAGCAAACATTTGAGTGGTTTTGACTCTAATCCAGGCCTGGT 1354 1397 ACTTGTCCCAGCAAACATTTGAGTGGTTTTTAATCCACGACCTTCTGATGAT 1354 1398 GTTAAACCTTTGCAAGAACATTGACCCACTGATTAAGTCCACCAGGTTTTTCCAGATGAT 1354 1399 GTTAAACCTTTGCAAGAACATCACCAGCTGATAAGTCCACCAGTTTTTCCAAGAAAT 1453 1355 GTCATCAGTTTCATAAAGCGGCACACTGTGATGATAAAGTCCCAGTTTCCTATGAACAAT 1453	1454 CGCCCARTAGTGATCAAAACGGATGTAAATTATCAATTTACACAAATTGCGTAGACCGA 1513 1415 GGACCAACGTTCAAGAGAATCAATGTGGATCACTACAAATTGCGAACGATGATGTGGTGGATCAT 1474 1514 GTGGATGCAGAAGATCAATGTGTATTGTTTATCGGAACAGATGTGGACCGTT 1573 1475 GTCATTGCAGAAGATCAATGTTATTTTTTTTTTTTTTTT
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TTAAGGTAACCCTGGAAGTCATTGACACAGAGCATTTGGAAGAACTTCTTCAT 2233 KAAAGCGTGTGCTGAGTGTTGCCTCGCCCGAGACCCTTACTGTGCTTGGGAT 1813 ATGITCICGCIATITICCCACIGCAAAGAGACGCACAAGACGACAAGAIAIA 1873 AGACCCACTGACTCACTGTTCAGACTTACACCATGATAATCACCATGGCCAC 1933 AGAGAGAATCATCTATGGTGTAGAGAATAGTAGCACATTTTTGGAATGCAGT 1993 COCAGAGAGCGCTGGTCTATTGGCAATTCCAGAGGCGAAATGAAGAGCGAAAA 2053 CAGAGTGGATGATCATCAGCACAGATCAAGGCCTTCTGCTACGTAGT 2113 GAAGCCCGATGAAAGAATCATCAAAACGGAATATGGGCTACTGATTCGAAGT 2068 GAAGGATTCAGGCAATTACCTCTGCCATGCGGTGGAACATGGGTTCATACAA 2173 TGATG 2246 TGAGG 2201

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/cgn2_6/ptodata/2/ina/PCTUS_COMB.seq:*
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GenCore version 5.1.6
Copyright (c) 1993 - 2004 Compugen Ltd.
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US-09-308-115-3

US-09-308-119B-2

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COMPUTE READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC COMPATION
COMPUTER: PALLICATION DATA:
APPLICATION NUMBER: US/08/121,713D
FILING DATE: 13-SEP-1993
CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
NAME: COMMAN RICHARD 36,627
REGISTRATION NUMBER: 36,627
REBERENCE/DOCKET NUMBER: 36,627
REBERENCE/COCKET NUMBER: 36,627
REBERENCE/COCKET NUMBER: 34,627
TELEPHONE: (415)343-4341
TELEFAK: (415)343-4341
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APPLICANT: Kolodkin, Alex L.
APPLICANT: Matthes, David
APPLICANT: Bentley, David R.
APPLICANT: O'Connor, Timothy
TITLE OF INVENTION: The Semaphorin Gene Family
NUMBER OF SEQUENCES: 100
CORRESPENDENCE ADDRESS:
ADDRESSEE: SCIENCE & TECHNOLOGY LAW GROUP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ALIGNMENTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 53, Application US/08121713D Patent No. 5639856
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SEQUENCE CHARACTERISTICS:
LENGTH: 2601 base pairs
                                                   LENGIR: ZOTE ACID
TYPE: nucleic acid
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Best Local Similarity
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CITY: San F1
STATE: CA
COUNTRY: US/
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US-08-121-713D-53
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Matches 2509; Conservative 0; Mismatches 1; Indels 0; Gaps 0; 193 CTGCAGCATGGGCTGGTTAACTAGGATTGTCTTTTTTGGGGAGTATTACTACAGC 252	TACCTTCCTTTTGGATGAGGAACGGAGTAGGCTGTATGTTGGAGCAAAGGATCACATATT TACCTTCCTTTTGGATGAGGAACGGAGTAGGCTGTATGTTGGAGCAAAGGATCACATATT TACCTTCCTTTTTGGATGAGGAACGGAGTAGGCTGTATGTTGGAGCAAAGGCAACGATCTTAGACTTTCATTCGACCTGGTTAATTTCAAAAGATTGTTGTGTGGCCAGTATCTTACACCTGAAGATTGTTGGACCTGGATATTTCATCTTACACCTGGTTAATTTCATGAGGATTTTCATGAGGATGATGTTGTGTGTG	TCCAATTICGACCTACATTGAAATTGGACATCCTGAGGACAATATTTTAAGCTGGA TCCAATTTGCACCTACATTGAAATTGGACATCATCCTGAGGACAATATTTTTAAGCTGGA TCCAATTTGCACCTACATTGAAATTGGACATCCTGAGGACAATATTTTTTAAGCTGGA GAACTCACATTTTGAAAAACGGCCGTGGGAAAGATCCATATGACCTGATGCTGGAGG GAACTCACATTTTAATAGAAGGACCGTGGGAAGATCCATATGACCCTAAGCTGCTGACGC ATCCCTTTTAATAGATGGAGAATTATACTCTGGAACTGCAGCTGATTTTATGGGGCGAGA ATCCCTTTTAATAGATGGAAATTATACTCTGGAACTGCAGCTGATTTTATGGGGCGAGA CTTTGCTATTTTAATAGATGGAAATTATACTCTGGAACTGCAGCTGATTTTATGGGGCGAGA CTTTGCTATCTTCCGAACTCTTGGGCACCACCCCAATCAGGACAGAGCAGCAATGTTC [S53 CAGGTGGCTCAATGATCAATGATCCACCCCACTCATCTCAGAGAGTGACAATCC 1
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92.6%; Score 2508.4;
Best Local Similarity 100.0%; Pred. No. 0;
Matches 2509; Conservative 0; Mismatches
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GGATGAGTTCTGTGAACAAGTTTGGAAAAGGGACCGAAAACAACGTCGGCAAAGGCCCAGG
                                                2169 GGATGACTTCTGTCAAGTTTGGAAAAGGGACCGAAACAACAACGTCGGCAAAGGCCAGG
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COMPUTE: 94104
COMPUTE: P10ppy disk
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Patent No. 5807826
GENERAL INFORMATION:
APPLICANT: Goodman, Corey S.
APPLICANT: Rolodkin, Alex L.
APPLICANT: Bentley, David APPLICANT: Bentley, David NAPLICANT: O'Connor, Timothy
TITLE OF INVENTION: The Semaphorin Gene Family
NUMBER OF SEQUENCES: 100
CORRESPONDENCE ADDRESS:
ADDRESSEE: SCIENCE & TECHNOLOGY LAW GROUP
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SEQUENCE CHARACTERISTICS:
LENGTH: 2601 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
TOPOLOGY: linear
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969 TCCTAAAAATCCAGTTGTATAGGAGTGTTTACGACTTCCAGTAACATTTCAAGGGATC 1028 1213 AGCCGTGTGTATGTATAGCATGAGTGTTTACGACTTCCAGTAACATTTTCAAGGGATC 1028 1029 AGCCGTGTGTATGTATAGCATGAGAGGTGTTCCTTGGTCCATATGCCCA 1088 1273 CAGGGATGACCCAACTAATCAATGGGTGTTCCTTGGTCCTTATGGCCCA 1088 1273 CAGGGATGGACCCAACTATCAATGGGTGCTTATCAAGGAAGAGTCCCCTATCCAGGGC 1332	TGCCCAATAGTAACATCATCCAGCCATGTACAATCCAGTGTTTCCTATGAACAA TGGCCCAATAGTGATCAAACGATGTAATTATCAATTTACACAATTGTCGTAGACCA TGGCCCAATAGTGATCAAACGGATGTAAATTATCAATTTACACAAATTGTCGTAGACCG TGGCCCCAATAGTGATCAAAACGGATGTAAATTATCAAATTGTCGTAGACCG AGTGGAAGGTGCAGAAGATGAAGATAATAATCAAAATTGTCGTAGACCG AGTGGAAGGTGCAGAAGATGGAAGATTAATCGAAATTTACACAAATTTGTCGTAGCCGT TGTTAAAGTAGAAGATGGACAGTATGATGTTATTATCGGAACAGATGTTGGGACCGT TCTTAAAGTAGTATCCTAAGGAAGACTTGGTATGTTTATAAAAGAGGTTCTGCTGGA	AGAAATGACAGTTTTTCGGGAACCGCTTTTCAGCAATGGGGGCTTTCCACTAAGCA	1689 AAGAATGGAGCCCACTGATCACTGTTCAGACTTACACCATGATAATCACCATGGCCX 1748 1933 CAGCCCTGAAGAGAATCATCTATGGTGTAGAGAATAGTAGCACATTTTTGGAATGCAG 1992 1749 CAGCCCTGAAGAGAATCATCTATGGTGTAGAATAGTAGCACATTTTTGGAATGCAG 1808 1993 TCCGAAGTCGCAGAGGGCTGGTTATTGGCAATTCCAGAGGCGAAATGAAGAGCGAAA 1868 2053 AGAAGAGATCGAGAGGCTGGTTATTGGCAATTCCAGAGGCGAAATGAAGAGCGAAA 1868 2053 AGAAGAGATCAGAGAGGCTGGTTATTGGCAATTCCAGAGGCGAAATGAAGAGCGAAA 1868 2053 AGAAGAGATCAGAGAGGCTCTATTAGCAATTCCAGAGGCGAAATGAAGAGCGAAA 1868 2113 TCTACAACAGAAGAGTGATCATATCATCAGACAATCAAGAGCTTCTGCTACGTAG 1928 2114 AGAAGAGATCAGAAGGAATTACCTCTGCCATGCGGTGGAACTTCTTCACA 1988 2173 AACTCTTCTTAAGGTAACCTGGAAATTACCTTGACATCGGTGGAACATGGATTCATCA 232

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Er: Floppy disk

IBW PC compatible
SYSTEM: PC-DOS/MS-DOS
PATENTIN Release #1.0, Version #1.25
ACTION DATE:

N NUMBER: US/09/060,692
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Kolodkin, Alex L.
Matthes, David R.
O'Connor, Timothy
O'Connor, The Semaphorin Gene Family
OUENCES: 100
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Bush Street, Suite 3200
rancisco
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1. UNBER: US/08/121,713
TI INFORMATION:
TO RICHARD 36,627
TO NUMBER: B94-002-1
COCKET NUMBER: B94-002-1
(415) 343-4341
(415) 343-4342
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ACTERISTICS:
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STRANDEDNESS: doul TOPOLOGY: linear MOLECULE TYPE: cDNA FEATURE:
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1989 AACTCTTCTTAAGGTAACCCTGGAAGTCATTGACACAGGGCATTTGGAAGAACTTCTTCA 2048
                                                                                                                                   CCAGAAGGTCTGGTACAGACTTCATGCAGCTCATCAACCACCAATCTCAACAGGT 2352
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Goodman, Corey S.
APPLICANT: Kolodkin, Alex L.
APPLICANT: Kolodkin, Alex L.
APPLICANT: Matthes, David
APPLICANT: O'Connor, Timothy
TITLE OF INVENTION: The Semaphorin Gene Family
CORRESPONDENCE: 100
CORRESPONDENCE ADDRESS:
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Patent No. 6013781
GENERAL INFORMATION:
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-
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TELEPHONE: (415)343-4341
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ATTORNEY/AGENT INFORMATION:
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy of
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: U
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               INFORMATION FOR SEQ ID NO: 5
SEQUENCE CHARACTERISTICS:
LENGTH: 2601 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                               Query Match
Best Local Similarity 100.0
Matches 2509; Conservative
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                                                                                                                    TOPOLOGY: linear MOLECULE TYPE: cDNA
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TCTGGTGAATAAATGGACAACATTCCTCAAAGCTCGTCTGATTTGCTCAGTGCCAGGTCC 1092
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REGISTRATION NUMBER: 36,627
REFRERNCY-DOCKET NUMBER: B94-002-1
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415)343-4341
                                                                                                                                    INFORWATION FOR SEQ ID NO: 53:
SEQUENCE CHARACTERISTICS:
LENGTH: 2601 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
                                                                                         TELEFAX: (415) 343-4342
                                                                                                                                                                                                                                                                                                                              NAME/KEY: CDS
LOCATION: 16..2331
                                                                                                                                                                                                                                                            TOPOLOGY: linear MOLECULE TYPE: CDNA.
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          (415) 781-1989
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Matches 2509, Conservative
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PCT-US94-10151A-53
       TELEPHONE:
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Sequence 53, Application PC/TUS9410151A
GENERAL INFORMATION:
APPLICANT: THE Regents of the University of California
TITLE OF INVENTION: The Semaphorin Gene Family
NUMBER OF SEQUENCES: 66
CORRESPONDENCE ADDRESS:
ADDRESSPEN: FLERH HOHBACH TEST ALBRITTON & HERBERT
STREET: 4 Embarcadero Center, Suite 3400
CITY: San Francisco
STREE: CAS STREET: USA
COUNTRY: USA
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CMPUTER REAABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: DATOR PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US94/10151A
FILING DATE: 13-SEP-1994
CLASSIFICATION:
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NAME: Osman, Richard A.
REGISTRATHON NUMBER: 36,627
REFERENCE/DOCKET NUMBER: FP-58750-PC/RAO
TELECOMMUNICATION INFORMATION:
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NAME: Deluca, Mark
REGISTATION NUMBER: 33,229
REFERENCE/DOCKET NUMBER: UFN-1428
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE:
CLASSIFICATION: 435
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                                      TCTGGTGAATAAATGGACAACATTCCTCAAAGGTCGTCTGATTTGCTCAGTGCCAGGTCC 1092
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                                                                            849 TCTGGTGAATAAATGGACAATTCCTCAAAGCTCGTCTGATTTGCTCAGTGCCAGGTCC 908
                                                                                                                                                 AAATGGCATTGACACTCATTTTGAAGGAACTGCAGGATGTATTCCTAATGAACTTTAAAGA 968
789 AAAAGCTACTCACGCTAGAATAGGTCAGATATGCAAGAATGACTTTGGAGGGCACAGAAG 848
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APPLICANT: Raper, Jonathan A.
APPLICANT: Luo, Yuling
TITLE OF INVENTION: Compositions Which Regulate Neural
TITLE OF INVENTION: Regeneration and Methods of Making and Using the Same
NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
ADDRESSEE: Woodcock Washburn Kurtz Mackiewicz &
ADDRESSEE: No. 5416197ris
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2472
1929 TCTACAACAGAAAGATTCAGGCAATTACCTCTGCCATGCGGTGGAACATGGGTTCATAAA 1988
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                                                                                                                                          1989 AACTCTTCTTAAGGTAACCCTGGAAGTCATTGACACAGAGGATTTGGAAGAACTTCTTCA
                                                                                                                                                                                                                                         TAAAGATGATGATGGAGATGGCTCTAAGACCAAAGAAATGTCCAAATAGCATGACACCTAG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2109 ccagaaggrcrggracagagacrrcargcagcrcarcaaccacccaarcrcaacacgar
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2469 CCAATTATAAATTAAAATCCATGAGTAACTTTCCTAATAGGCTTTTTTTC 2518
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/136,922
```

TELECOMMUNICATION INFORMATION:
TELEPHONE: 215-568-3100
TELEFAX: 215-568-319
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 1481 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
TOPOLOGY: both
MOLECULE TYPE: cDNA
FEATURE:), NAME/KEY: CDS ; LOCATION: 50..1480 US-08-136-922-1

TUR Seb 73 00:34:0/ 7004

953 TACTGTGCTTGGGATGGTTCTGCATGTTCTCGCTATTTTCCCACTGCAAAGAGACGCACA 1012

1799 TACTGTGCGATGGTTCTGCATGTTCTCGCTATTTTCCCACTGCAAGAGACGCACA

TTACACCGGTGTGTATTTACGGGAAAGCGTGTGCTGAGTGTTGCCTCGCCCGAGACCCT

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CTTTCCACTAAGCAGCAACAACTATATATTGGTTCAACGGCTGGGGTTGCCCAGCTCCCT TYACACCGGTGTGATATTTACGGGAAAGCGTGTGCTGAGTGTTGCCTCGCCCGAGACCCT 1859 AGACGACAAGATATAAGAAATGGAGACCCACTGACTCACTGTTCAGACTTACACCATGAT 1073 AATCACCATGGCCACAGCCCTGAAGAGAGAAATCATCTATGGTGTAGAGAAATAGTAGCACA

AATCACCATGGCCACAGGCCCTGAAGAGAATCATCTATGGTGTAGAGAATAGTAGCACA

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1138 1378 1438 1018 1078 ATTITCAAGGGATCAGCCGTGTGTATGTATAGCATGAGTGATGTGAGAAGGGTGTTCCTT 1258 GGTCCATATGCCCACAGGGATGGACCCAACTATGATGGGTGCCTTATCAAGGAAGAGTC 1318 ATGAACTTTAAAGATCCTAAAATCCAGTTGTATATGGAGTGTTTACGACTTCCAGTAAC 1198 532 172 CCCTATCCACGGCCAGGAACTTGTCCCCAGCAAAACATTTGGTGGTTTTTGACTCCTACAAAG cecharecaegecaegaacrigicecaecaaaacarrigeregrifitaacretacaaag GACCITICCTGATGATGTTATAACCITTGCAAGAAGTCATCCAGCCATGTACAATCCAGTG 53 GAACATGACAATCCTGAAGAAGACAAAGTATACTTTTTCTTCCGTGAAAATGCAATAGAT 959 GGAGAACACTCTGGAAAAGCTACTCACGCTAGAATAGGTCAGATATGCAAGAATGACTTT 113 GGAGAACACTCTGGAAAAGCTACTCACGCTAGAATAGGTCAGATATGCAAGAATGACTTT TCAGTGCCAGGTCCAAATGGCATTGACACTCATTTTGATGAACTGCAGGATGTATTCCTA 899 GAGAGTGACAATCCTGAAGATGACAAAGTATACTTTTTTCTTCCGTGAAAATGCAATAGAT 1019 GGAGGGCACAGAAGTCTGGTGAATAAATGGACAACATTCCTCAAAAGCTCGTCTGATTTGC Gaps 0; Length 8; Indels 1; DB ; Score 1415.2; ; Pred. No. 0; 0; Mismatches 52.2%; Similarity 99.4%; %0; Conservative 0 Query Match Best Local Simil Matches 1420; (1079 1139 1199 353 1259 413 1319 1379 293 473 B ò 중 음 d qq 8 8 δ g 셤 Q $\stackrel{>}{\circ}$ δ

1253 CTTCTGCTACGTAGTCTACAACAGAAGGATTCAGGCAATTACCTCTGCCATGCGGTGGAA 1312

CTTCTGCTACGTAGTCTACAACAGAATTCAGGCAATTACCTCTGCCATGCGGTGGAA

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1193 AATGAAGAGGGAAAAAAAAAAGAATCAGGATCATCATATCAGGACAGATCAAGGC

2039 ANTGAAGAGCGAAAAGAAGAGATCAGAGTGGATGATCATATCATCAGGACAGATCAAGGC

CATGGGTTCATACAACTCTTCTTAAGGTAACCCTGGAAGTCATTGACAAGGGAGTTG GAAGAACTTCTTCATAAAGATGATGATGGAGATGGCTCTAAGACAAAGAAATGTCCAAT

CATGGGTTCATACAAACTCTTAAGGTAACCTGGAAGTCATTGACACAGAGCATTTG

Sequence 2, Application US/09308179B
Patent No. 643666
GENERAL INFORMATION:
APPLICANT: INAGARI, Shinobu
APPLICANT: INAGARI, Shinobu
TITLE OF INVENTION: NOVEL SEMAPHORIN GENES (I)
FILE REFERENCE: 0020-4562P
CURRENT FILING DATE: 1999-05-14
PRIOR APPLICATION NUMBER: US/09/308,179B
CURRENT FILING DATE: 1999-11-15
PRIOR APPLICATION NUMBER: PCT/JP97/04111
PRIOR APPLICATION NUMBER: DAPAN 321068/1996
PRIOR APPLICATION NUMBER: JAPAN 321068/1996
NUMBER OF SEQ ID NOS: 4
SOFTWARE: PatentIn Ver: 2.1
SEQ ID NO 2
LENGTE: 2898 Ĥ TYPE: DNA ORGANISM: Mus sp. RESULT 8 US-09-308-179B-2 FEATURE:

t, unknown, or NAME/KEY: misc_feature LOCATION: (1)...(2898) GOTHER INFORMATION: Strandedness: Double-stranded FEATURE: NAME/KEY: misc_feature LOCATION: (1)...(2898) GOTHER INFORMATION: any n = a, c, g, t, unknown, c US-09-308-179B-2

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1499 ATTGTCGTAGACCGAGTGCATGCAGAAGATGGACAGTATGATGTTTATCGGAACA 1558

1559 GATGTTGGGACCGTTCTTAAAGTAGTTTCAATTCCTAAGGAGACTTGGTATGATTTAGAA

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TITCCIATGAACAATCGCCCCAATAGTGATCAAAACGGATGTAAATTATCAATTTACACAA 1498

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CTTTCCACTAAGCAGCAACAACTATATTGGTTCAACGGCTGGGGTTGCCCAGCTCCCT 1738

1619 773 1679

Query Best Match	Query Match 22.0%; Score 596.4; DB 4; Length 2898; Best Local Similarity 56.3%; Pred. No. 4.5e-176; Matches 1229; Conservative 0; Mismatches 931; Indels 24; Gaps 5;		TGTGAGAGGGGTGTTCCTTGGTCCATATGCCCACAGGGATGGACCCAACTATCAATGGGT 129
. yo da	223 CTGTCTTTTCTGGGGGGTATTACTTACAGGGGAAACTATCAGAATGGGAAGAACAA 282 396 CTTGCTGGTCTGGGGTCACCTGCTGGAACTCTGGACCCCAGGTCACTCCGCGAACCCCTC 455		GCCITATCAAGGAAGACCCCIATCCACGGCCAGGAACTIGTCCCCCAAACATTGG 133 ACTATATGAAGGAAAAGTCCCCTACCCAAGGCCTGGTTCCTGTGCCCAGGAAAGTAAACGG 153
λ 6	283 IGTGCCAAGGCTGAAATTATCCTACAAAAAAAAAATGTTGGAATCCAATGTGAATCACTTT 342 		TGGTTTTGACTCTACAAGGCCTTCCTGATGATGTTATAACCTTTGCAAGAGGCCAAGAGGCCATGCGGTTGGCGGC
<u> </u>	3 CAATGGCTTGGCCAACAGCTCCAGTTATCATACCTTCCTT		1417 TCCAGCATGTACAATCCAGTGTTTCCTATGAACAATCGCCCAATAGTGATCAACGAA 1476
1 & 1	GCTGTATGTTGGAGCAAAGGATCACATATTTTCATTCGACCTGGTTAATATCAAGGA	Å qq	1477 TGTAAATTATCAATTTACACAAATTGTCGTAGACCGGGTGGGATGCAGAAGAGGGAGG
8 8 8	GGCTGG		1537 TGATGITATGATITATGGGAAGAIGTTGGGACCGTTCTTAAAGTAGTTTCAATTCCTAA 1596 1713 TGACGICTTATTATGGGACAGACACAGGAATTGTGCTGAAAGTAATCACAATTTACAA 1772
3 <i>i</i> 5	AAAAGACATCCTGAAAGAATGTGCTAATTTCATCAAGGTACTTAAGGCATATAATCAGAC	& a	1597 GGAGACTIGGIAIGÀITIAGAAGAGGITCIGCIGGAAGAAATGACAGITITICGGGAACC 1656 1773 CCAAGAACAGAGTGGAGGAGGAAGTCAITCTAGAGGAACTICAAAIAITCAAGGATCC 1832
සි දි	696 AAAAGACGCAAATGAGTGCCAATTATATCCGGGTTTTGCATCACTACAACAGGAC 752 580 TCACTTGTACGCCTGTGGAACGGGGGCTTTTCATCCAATTTGCACCTACATTGAAATTGG 639	÷ 8	1657 GACTGCTATTTCAGCAATGGAGCTTTCCACTAAGCAGCAACTATATATTGGTTCAAC 1716
9a &	753 ACACCTTCTCACCTGTGCTACTGGAGCTTTTGATCCACACTGTGCCTTCATCAGAGTCGG 812 640 ACATCATCCTGAGGACAATATTTTAAGCTGGAAACTCACATTTTGAAAACGGCGTGG 699	٠	
8 & 8 B	813 GCACCATTCAGAGGAACCCCTGTTTCACTGGAGTCACACAGAATCTGAGAGAGGGG 872 700 GAAGAGTCCATATGACCCTAAGCTGCTGACAGCATCCCTTTTAATAGAGGAATGAAT		GIGITGCCTCGCCGAGACCCTTACTGTGCTTGGGATGGTTCTGCATGTTTT 1
q ₀	873 CAGATGTCCTTTTGACCCCCAACTCCTCTTGTGTCCACGCTAGTTGGGAATGAGCTGT 932	셤 &	1953 CTGCTGCTGGCTCGAGACCCGTACTGTGCCTGGGATGGCATATCCTGCTCCAGGTACTA 2012 1837 TCCCACTGCAAAGAGACGCACAAGACGACAAGATATAAGAATGGAGACCC 1887
දුරු දුර	760 CTCTGGAACTGCAGCTGATTTTATGGGGGGGGAGACTTTGĆTATCTTCCGAACTCTTGGGCA 819 		
8 8 1	CCACCACCAATCAGGACAGAGCAGGATGATTCCAGGTGGCTCAATGATCCAAAGTTCAT	, v	1888 ACTGACTCACTGTTCAGACTTACACCATGATAATCACCATGGCCACAGCCCTGAAGAGAG 1947 10
g &	993 GITAGGCCATAITCGCACTGAGCATGAGGATGGGCTCCTGAAAGAACCAAAAITTGI 1052 880 TAGTGCCCACCTCAGAGAGAGAGAAACTAAAGAAAGAAAG	8 8	1948 AATCATCTATGGTGTAGAGAATAGTAGCACATTTTTGGAATGCAGTCCGAAGTCGCAGAG 2007
dg vo	1053 AGGTTCATATATGATTCCTGATAACGAAGACCGAGATGACAACAAATGTACTTTTTCTT 1112 940 CCGTGAAAATGCAATAGATGGAGAACACTTGGAAAAGCTACTCACGCTAGAATAGGTCA 999		AGCGCTGGTCTATTGGCAATTCCAGAGGCGAAATGAAGAGCGAAAAGAAGAGGATCAGAGT
2 d		음 &	2193 ÄGCAAAAGTCATCTGGTTTGTACAGAAGGGACGCGACGTAAGAAAAGAAGGAGGGAG
۲۵ Ag	1000 GATATGCAAGAATGACTTTGGAGGCACAGAAGTCTGGTGAATAATGGACAACATTCCT 1059		
} &	CAAAGCTCGTCTGATTTGCTCAGTGCCAAATGGCATTGACACTCATTTTGATGA	 & 8	2128 TTCAGGCAATTACCTCTGCCATGCGGTGGAACATGGGTTCATACAAACTCTTCTTAAGGT 2187 2313 TGCAGGGACCTATTTTTGCCAGACAGTAGAACACAATTTTGTCCATACTGTGCGTAAAAT 2372
음 장	1233 TAAAGCGCGGCTGGTTTGCTCAGTGCCGGGAATGAATGGAATCGACATACTTTGACGA 1292 1120 ACTGCAGGATGTATTCCTAATGAACTTTTAAAGATCCTAAAAATCCAGTTGTATATGGAGT 1179	8 8	2188 AACCCTGGAAGTCATTGACACAGAGCATTTGGAAGAACTTCTTCATAAAGATGATGATG- 2246
Db	ACTAGAGGATGTGTTTTTACTGCCGACCAGAGATCCTAAGAATCCAGTGATATTTGGACT		GAGATGGCTCTAAGACCAAAGAAAGAAAGTCTCCAATAGCATGACGTCTAGCCAGAAGGT
Oy Dp	1180 GTTTACGACTTCCAGTAACATTTTCAAGGGATCAGCCGTGTGTATAGTATAGATGATGAGTGA 1239	_	2433 GGAAAGACATCACAAGAIGCCCTGCCCTCCCTTAAGCGGTAIGTCTCAGGGGACAAACC 2492
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1460 ATAGTGATCAAAACGGATGTAAATTATCAATTTACACAAATTGTCGTAGACCGAGTGGAT 1519 1520 GCAGAAGAIGGACAGIAIGITIAIGITIAICGGAACAGAIGITGGGACCGIICIIAAA 1579 1580 GIAGITICAAITCCIAAGGAGACITGGIAIGATITAGAAGAGGITCTGCTGGAAGAAATG 1639 1628 TİTTİGGÜCAGAATAGGAAATAĞIĞÜTİTİTÜTAAATĞACAĞCCİTITİCCİĞĞAĞĞAĞAİĞ 1687 1388 GCTGGCTCATCCTCCTTAGAAAGATATGCAACCTCCAATGAGTTCCCTGATGATACCCTG 1447 1448 AACTICATCAAGACGCACCCGCTCATGGATGAGGCAGTGCCCTCCATCTTCAACAGGCCA 1507 1508 IGGTİCCİGAĞAÇAATGĞICAĞATACĞCCITAĞCAAATTĞCAĞTĞĞAĞĞÇAÇAĞĞTĞCI 1567 1568 GGGCATATCAGAATCACACTGTGGTTTTTCTGGGATCAGAGAAGGGAATCATCTTGAAG 1627 1268 TATGACATGCTTGCCAGTGTTTTTACTGGGACATTCAAGGACAACAGAAGTCTCT 1327 765 GAACTGCAGCTGATTTTATGGGCCGAGACTTTGCTATCTTCCGAACTCTTGGGCACGCC 824 1343 CCCAGCAAA---ACATTTGGTGGTTTTGACTCTACAAAGGACCTTCCTGATGATGTTATA AACTATCAATGGGTGCC----TTATCAAGGAAGAGTCCCCTATCCACGGCCAGGAACTTGT 1328 garriccaccriegacacceaerriccrigargaacgagirriccraagcccaggccaggrigir 1400 ACCITIGCAAGAAGICAICCAGCCAIGIACAAICCAGIGIIIICCIAIGAACAAICGCCCA Sequence 1, Application US/09077940A

Sequence 1, Application US/09077940A

Sequence 1, Application US/09077940A

GENERAL INFORMATION:

APPLICANT: KIMUZA, Toru et al.

TITLE OF INVENTION: NOVEL SEMAPHORIN Z AND GENE ENCODING THE SAME

TITLE OF INVENTION: NOVELS: 1998-06-05

CURRENT FILING DATE: 1998-06-05

NUMBER OF SEQ ID NOS: 20

SOFTWARE: Patentin version 3.1

LENGTH: 3692 33; Length 3692; 3.5%; Score 95.2; DB 4; Length 3 47.8%; Pred. No. 6.7e-19; tive 0; Mismatches 438; Indels TYPE: DNA ORGANISM: Rattus norvegicus 3.55 Query Match Best Local Similarity 47.85 Matches 431; Conservative ; LOCATION: (2683)..(3653) ; CTHER INFORMATION: ; NAME/KEY: POLYA.site ; LOCATION: (3654)..(3692) ; OTHER INFORMATION: US-09-077-940A-1 FEATURE:
NAME/KEY: 5'UTR
LOCATION: (1)..(18)
OTHER INFORMATION:
NAME/KEY: CDS
LOCATION: (19)..(2682)
OTHER INFORMATION:
NAME/KEY: 3'UTR
LOCATION: (2683) 1640 ACAGTTT 1646 1688 AGTGTTT 1694 RESULT 10 US-09-077-940A-1 1286 g à δ ò g ò В g 8 d g ò ઠે ð ò ద Sequence 1002, Application US/09976594

Better No. 6673549

GENERAL INFORMATION:
APPLICANT: Furness, Michael
APPLICANT: Furness, Michael
APPLICANT: Buchbinder, Jenny
TITLE OF INVENTION: GENES EXPRESSED IN C3A LIVER CELL CULTURES TREATED WITH STEROIDS
FILE REPRENCE: PA-0041 US
CURRENT APPLICATION NUMBER: US/09/976,594
CURRENT FILING DATE: 2001-10-12
NUMBER OF SEQ ID NOS: 1143
SOGTWARE: PERL PROGram
SEQ ID NO 1002
LIGHTH: 2278 GITGIATATGGAGTGITTACGACTTCCAGTAACATTTTCAAGGGATCAGCCGTGTATG 1225 1046 TGGACAACATTCCTCAAAGCTCGTCTGATTTGCTCAGTGCCAGGTCCAAATGGCATTGAC 1105 1097 recacercerrechea a decede contra a contract recorde ----- a da crement 1150 1106 ACTCATTTTGATGAACTGCAGGATGTATTCCTAATGAACTTTAAAGATCCTAAAATCCA 1165 1151 TTTTATTTCAACATTCTCCAGGCAGTTACAGATGTGATTCGTATCAACGGGGGGTGAT-- 1207 1226 TATAGCATGAGTGATGTGAGAAGGGTGTTCCTTGGTCCATATGCCCACAGGGATGGACCC 1285 989 AGAATAGGTCAGATATGCAAGAATGACTTTGGAGG---GCACAGAAGTCTGGTGAATAAA 1045 1037 AGAGTGGCTTCAAAAAAAAAAAAAAAAAAAAAAAAGAGTCCTGGAGAAACAG 1096 Grigicergecaacerricadacerraraadacaareergeerergearergicee 1267 9 GTGGTACAAGGAATTCTTGCAGCTGATTGGCTACAGCAACTTCCAGAAGTGGAAGAATA 2552 938 ccaracrirgricaagccgraga--------riracgagariaric 976 TACTITITICITICGIGAAAATGCAATAGATGGAGAACACTCTGGAAAAGCTACTCACGCT 988 817 808 868 Agrichiegagaaagiccnacccigcaaccicchaagacairicaaanigchigaaaaa 937 CCAAAGTICATTAGTGCCCACCTCATCTCAGAGAGTGACAATCCTGAAGATGACAAAGTA 928 689 AACGGCCGTGGGAAGAGTCCATATGACCTAAGCTGCTGACAGCATCCCTTTTAATAGAT 748 877 758 adcidosartodecidariandartodecida de accentoda de accentoda de accentoda de accentoda de accento GGAGAATTATACTCTGGAACTGCAGCTGATTTTATGGGGCGAGACTTTGCTATCTTCCGA egaaaacriaracricaeccacacaereacreacricerrecearreaceaerearracee ACTETTGGGCACCACCCAATCAGGACAGACAGCATGATTCCAGGTGGCTCAATGAT 977 TACTICITCTICAGGAAATAGCAGTGGAGTATAACACCATGGGAAAGGTAGTTTTCCCA Gaps 39; Query Match

4.6%; Score 125; DB 4; Length 2278;
Best Local Similarity 50.5%; Pred. No. 2.2e-28;
Matches 488; Conservative 0; Mismatches 440; Indels 38 ; NAME/KEY: misc_feature ; OTHER INFORMATION: Incyte ID No. 6673549 411373.7 US-09-976-594-1002 2385 2553 Criccananacircricitatacaca 2576 CTGTGAACAAGTTTGGAAAAGGGA TYPE: DNA ORGANISM: Homo sapiens 1166 1208 818 929 878 869 749 809 2493 2362 Dp 쉱 ò g ₽ d à 셤 õ d ò qq ð ద ò δ ò

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49.0%; Pred. No. 1.2e-17;
cive 0; Mismatches 424; Indels
FILE REFERENCE: HYS-23
CURRENT APPLICATION NUMBER: US/09/653,274
CURRENT FILING DATE: 2000-08-31
PRIOR APPLICATION NUMBER: 09/491,404
PRIOR FILING DATE: 2000-01-10
NUMBER OF ESQ ID NOS: 13
SOFTWARE: PatentIN Ver. 2.1
SEQ ID NO 12
LENGTH: 1923
TYPE: DNA
ORGANISM: Homo sapiens
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Matches 444; Conservative
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; LOCATION: (1)..(1923)
US-09-653-274-12
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        502 CCACAGTAACTGACTTCCTAGCCATCGACGCTGTTATCTACCGTAGCCTTGGGGACCGGC 661
                                                                                                                       562 CCACACTGCGCACAGTAAAGCATGACTCCAAGTGGTTTTAAAGAGCCATACTTTGTGCATG 721
                                                                                                                                                                                                                 825 ACCCAATCAGGACAGAGCAGCATGATTCCAGGTGGCTCAATGATCCAAAGTTCATTAGTG
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Patent No. 6635742

Patent No. 6635742

Patent No. 6635742

APPLICANT: NFORMATION:
APPLICANT: Arerburn, Matthew C
APPLICANT: Arerburn, Matthew C
APPLICANT: Tang, Y. Tom
APPLICANT: Liu, Chenghua
APPLICANT: Liu, Chenghua
APPLICANT: Liu, Chenghua
APPLICANT: Dianac, Radoje T
TITLE OF INVENTION: Methods and Maaterials Relating to Semaphorin-Like
TITLE OF INVENTION: Polypeptides and Polymucleotides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAGCTCGTCTGATTTGCTCAGTGCCAAGGTCCAAATGGCATTGACACTCATTTTGATGAAC
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APPLICANT: Yeurg, Gorge Y
APPLICANT: Atterburn, Matthew C
APPLICANT: Atterburn, Matthew C
APPLICANT: Tang, Y. Tom
APPLICANT: Tang, Y. Tom
APPLICANT: Tang, Y. Tom
APPLICANT: Dranac, Radoje T
ITLE OF INVENTION: Methods and Maaterials Relating to Semaphorin-Like
ITLE OF INVENTION: Methods and Polypucleotides
CURRENT APPLICATION NUMBER: US/09/653,274
PRIOR TELING DATE: 2000-08-31
PRIOR FILING DATE: 2000-01-10
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Mize, Nancy K
Tang, Y. Tom
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APPLICANT: Boyle, Bryan J
APPLICANT: Arterbur, Matthew C
APPLICANT: Mize, Nancy K
APPLICANT: Mize, Nancy K
APPLICANT: Tang, Y. Tom
APPLICANT: Liu, Chenghua
APPLICANT: Drmanac, Radoje T
TITLE OF INVENTION: Methods and Maaterials Relating to Semaphorin-Like
TITLE OF INVENTION: Polypeptides and Polynucleotides
FILE REFERENCE: HYS.-21ypeptides
                                                                                                         1523 GAAGATGGACAGTATGATGTTATGTTTATCGGAACAGATGTTGGGACCGTTCTTAAAGTA
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CURRENT FILING DATE: 2000-08-31
PRIOR APPLICATION NUMBER: 09/491,404
PRIOR FILING DATE: 2000-01-10
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Patent No. 6635742
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GENERAL INPORMATION:
APPLICANT: McClelland, Michael
APPLICANT: Welsh, John APPLICANT: Trenkle, Thomas
TITLE OF INVENTION: Reduced Complexity Nucleic Acid Targets and Methods of ITLE OF INVENTION: Using Same
FILE REFERENCE: P-PH 3457
CURRENT APPLICATION NUMBER: US/09/300,958A
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PRICA APPLICATION NUMBER: 60/083,331
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                                                                                                               APPLICANT: HALL, Kathryn T.
APPLICANT: FREEMAN, Gordon J.
APPLICANT: SCHUTZE, Joachim L.
APPLICANT: BOUSSIOTIS, Vassiliki
APPLICANT: BOUSSIOTIS, Vassiliki
TITLE OF INVENTION: NUCLBIC ACIDS ENCODING CD100 MOLECULES
FILE REFERENCE: DFN-005CPA2
CURRENT APPLICATION NUMBER: US/08/556,422A
CURRENT FILING DATE: 1995-11-09
NUMBER: OF SEQ ID NOS: 7
SOFTWARE: FastSEQ for Windows Version 4.0
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                            Sequence 1, Application US/08556422A
Patent No. 6576754
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Best Local Similarity 49.0%;
Matches 619; Conservative
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LOCATION: (88)...(2673)
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ORGANISM: Homo sapiens
                                                                                            GENERAL INFORMATION:
      US-08-556-422A-1
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